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   WPI; 2004-562147/54
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                                                                                                                     21-JAN-2004; 2004WO-US001468
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           analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are useful for promoting motor neuron viability and axon degeneration, target muscle reinmervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence represents a MNTF motoneuronotrophic factor peptide analogue.
                                                                                                                                                                                     New motoneuronotrophic factor peptide analogue enhancing the viability motor neurons, useful for target muscle reinnervation, treating peripheral nerve injuries or neurodegenerative diseases and in wound
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to motoneuronotrophic factor peptide analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are useful for promoting motor neuron viability and axon degeneration, target muscle reinnervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence represents FSRYAR 6 mer domain motoneuronotrophic factor peptide
                                                                                                                                         Claim 7;
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Neuropeptide-Agonist; axon degeneration; muscle reinnervation;
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                                                                                                         present invention relates to motoneuronotrophic factor peptide
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Pred. No. 2.1e+06;
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peripheral nerve; neur
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ADQ90200 standard;
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                                                                                                                                                                                                                                    The present invention relates to motoneuronotrophic factor peptide analogue, where the peptide analogue enlances the viability of mot
                                                                                                                                                                                                                                                                          Claim 7;
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Neuropeptide-Agonist; axon degeneration; muscle reinnervation;
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peptide; 13
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21-JAN-2003;
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                                                                                                                                                                                                                  Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 6; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FSRYAR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nt invention relates to motoneuronotrophic factor peptide where the peptide analogue enhances the viability of motor the methods and compositions of the present invention are under the methods and compositions of the present invention are under the methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ko 1D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       _(first entry)
2003US-0441772P
                                                                                                                                                                                                                                                                           neurodegenerative; wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               motoneuronotrophic factor peptide analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 AA
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Pred. No. 4.2;
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are us for promoting motor neuron viability and axon degeneration, target my reinnervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence represents a MNTF motoneuronotrophic factor peptide analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                               The invention relates to an isolated polynucleotide consisting of motoneuronotropic factor (MUTF) -associated nucleic acid sequence comprising ARA17647 or ARA17648, its fragment or complement. Also described are the following: a composition comprising a first and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Motoneuronotropic factor (MNTF)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New motoneuronotrophic factor peptide analogue enhancing the viability motor neurons, useful for target muscle reinnervation, treating peripheral nerve injuries or neurodegenerative diseases and in wound
                                                                                                                                                                                                                                                             New isolated polynucleotide consisting of a motoneuronotropic factor (MNTF)-associated nucleic acid sequence, useful for promoting the survival, growth, proliferation and maintenance of mammalian neurons
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-386358/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-NOV-2003; 2003US-0518581P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JUL-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEA17675 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to motoneuronotrophic factor peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-562147/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            motoneuronotropic factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENE-) GENERVON BIOPHARMACEUTICALS LLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERVON BIOPHARMACEUTICALS LLC
                                                                                                                                                                                                  SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ∂
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                                                                                                                                                                                              ID NO 29; 34pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MNTF; cell growth; cell proliferation;
16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:29
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RESULT 7
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Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Motorneuronotrophic factor; MNTF-1; MNTF1-F6; human; axon regeneration; motoneuron; diagnose; treatment; disease; wound healing; scar tissue;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human MNTF1-F6
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                                             This sequence, represents a fragment of a novel human motoneurotrophic factor, MNTF1-F6. Such factors are used to promote regeneration of the axon of a motoneurone, to diagnose and treat motoneurone disease in a mammal or to accelerate wound healing whilst concomitantly minimising inhibiting scar tissue and/or keloid formation in an area associated with the control of the contro
                                                                                                                                                                                                                                                                                                                                                                                        Motoneurotrophic factor MNTF1-F3 and MNTF1-F6 - regeneration, diagnosing or treating motoneuron wound healing without scar formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-SEP-1996;
15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
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                                                                                                                                                                                                                                                                                                                   Claim 4; Fig 2B; 78pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-230703/20.
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96US-00751225.
97US-00928862.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6
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    axonal regeneration, the
                                                                                                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Pred. No.
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    polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for motoneuron
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Query Match Best Local Matches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peripheral nerve injury; musculoskeletal disorder; spinal cord injury; head injury; stroke; neuromuscular degenerative disease; nootropic; amyotrophic lateral sclerosis; peripheral neuropathy; neuroprotective; muscular dystrophy; AIDS; spinal muscular atrophy; multiple sclerosis; scar; myasthenia gravis; sensory neuronal function disorder; vulnerary
                             injuries, head injuries, strokes, neuromuscular degenerative diseases, amyotrophic lateral sclerosis, spinal muscular atrophy, peripheral neuropathy, diabetic peripheral neuropathy, peripheral neuropathy resulting from AIDS or radiation therapy for cancer, multiple sclerosis, muscular dystrophy, inhibition of scar formation, myasthenia gravis and sensory neuronal function disorders. The present sequence is human MNTF1-F6 protein used to illustrate the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     administered at a concentration of 5 ng-50 mg, whereas for inhibiting hereditary motoneurone disease, the dosage is 5-100 (especially 30-50)ng
                                                                                                                                        The invention relates to a method for promoting the survival, growth, proliferation or maintenance of mammalian neurons by administering motoneuronotrophic factors (MNTF). The method is useful for treating peripheral nerve injuries, musculoskeletal disorders, spinal cord
                                                                                                                                                                                                                                                 Promoting the survival, growth, proliferation or maintenance of neurons by administering motoneuronotrophic factors, useful for musculoskeletal and neurodegenerative disorders and spinal cord
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human motoneuronotrophic factor (MNTF)1-F6 protein.
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                                                                                                                                                                                                                     Claim 1; Fig 2B;
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les 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              motoneuronotrophic factor; MNTF; therapy; radiation therapy; eral nerve injury; musculoskeletal disorder; spinal cord injury;
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                                                                                                                                                                                                                      90pp; English
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human MNTF1-
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treating
injuries.
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RESULT 10
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                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                        The present invention relates to motoneuronotrophic factor peptide analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are useful for promoting motor neuron viability and axon degeneration, target muscle reinnervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence represents MNTF1-F6 33 mer motoneuronotrophic factor peptide analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chau
 SAPHO syndrome;
                     Propionibacterium acnes immunogenic protein #18652.
                                               27-FEB-2002
                                                                                          AAU57756 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                      Claim 2; SEQ ID NO 1; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         New motoneuronotrophic factor peptide analogue enhancing the viability motor neurons, useful for target muscle reinnervation, treating peripheral nerve injuries or neurodegenerative diseases and in wound
                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-562147/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peripheral nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           motoneuronotrophic; motor neuron; Nootropic; Neuroprotective; Vulnerary;
Neuropeptide-Agonist; axon degeneration; muscle reinnervation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MNTF1-F6 33 mer motoneuronotrophic factor peptide analogue
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synovitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neurodegenerative; wound healing.
                                              entry)
                                                                                                                                                                                                               100.0%;
 acne;
                                                                                             81
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                                                                                                                                                                                                               Score 31; DB
Pred. No. 11;
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pustulosis; hypertosis;
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osteomyelitis;
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                                                                                                                                                                   RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                     the treatment, prevention and diagnosis of medical conditions caused by the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO Syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central concrous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The college polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (ELISA). Note: The sequence data for this patent did not form part of the printed specification, but was contacted in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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02-JUN-2000;
07-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequences AAU39105-AAU68017 represent Propionibacterium acnes polypeptides. The proteins and their associated DNA sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Propionibacterium acnes vaccinating against and treating acne vulgaris.
                            Propionibacterium acnes predicted ORF-encoded polypeptide #18951.
                                                                 20-OCT-2003
                                                                                                                                 ABM54275 standard; protein; 81
                                                                                                                                                                                                                                                                                                                                                           Sequence 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 18951; 1069pp; English
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                                                                                                                                                                                                                                                                                                                                                                                         ftp.wipo.int/pub/published_pct_sequences
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6; Conserv
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re J, Zhang
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2000US-0216747P.
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                                                                (first entry)
                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptides and nucleic acids useful for diagnosing infections, especially useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mitcham JL, Wang S
, Jen S, Carter D;
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Pred. No. 28;
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Acne vulgaris;

antiseborrhoeic;

dermatological; antibacterial;

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RESULT 12
AAU57258
ID AAU57
XX
AC AAU57
XX
AC AAU57
XX
DT 27-FE
                                                                                                                                                                                                                                                                                                                                                                                              CC encoding a Propionibacterium acnes protein. The invention also relates to CC polypeptides encoded by the polynucleotides (ABM5524-ABM64336) and to CC immunogenic fragments of P. acnes polypeptides. The invention CC additionally encompasses expression vectors and host cells comprising a polypeptide of the invention; fusion proteins comprising a polypeptide of the invention; a comprising polypeptides of the invention; a comprising polypeptides of the invention; a comprising polypeptides of the invention; a composition comprising P. acnes polypeptides, comprising P. acnes polypeptides, comprising P. acnes polypeptides, composition (comprising P. acnes polypeptides, composition (comprising P. acnes polypeptides, companies polypeptides, composition (comprising P. acnes polypeptides, companies polypeptides, companies and kit companies and a method for inhibiting the development of P. acnes in a companies of the p. acnes polypeptides, polypucleotides, antibodies, fusion proteins, T cell populations or antigen-presenting cells that express the complexity populations or antigen-presenting or treating acne composition. The polymucleotides can also be used as probes or primers for composition of an immune response specific for a P. acnes for conclete acid hybridisation. The vaccine composition is useful for the composition is useful for the companies of the primers for companies and companies of the primers for companies and companies of the present contained within the P. acnes polymucleotides of the companies of the primers for the primers of the primers for the primers of the primers of the primers of the primers of the companies and polypeptide predicted to be encoded by an ORF (open companies). The present of the primers of the primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
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Zhang Y, Wang S, Jen S, I
Barth B, Vallieve-Douglass
                                          AAU57258;
                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 18951; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-381789/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-OCT-2001; 2001US-00978825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Propionibacterium acnes
  27-FEB-2002
                                                                                     AAU57258 standard; protein; 112 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunostimulant; immune response; vaccine.
                                                                                                                                                                                                                                                                                Local Similarity
les 6; Conserv
                                                                                                                                                                                               14
                                                                                                                                                                                                                                     1 FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORIXA CORP
                                                                                                                                                                                               FSRYAR 19
                                                                                                                                                                                                                                                                                                                                                                   81
                                                                                                                                                                                                                                                                                                                                                                   A,
                                                                                                                                                                                                                                                                                  Conservative
  (first entry)
                                                                                                                                                                                                                                                                                100.0%; Score 31; DB 6; Length 81; 100.0%; Pred. No. 28; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Persing DH,
Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bhatia A,
Benson DR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Maisonneuve JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Carter D;
                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                Gaps
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  RESULT 13
ABM53777
ID ABM53
XX
AC ABM53
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AC ABM53
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    20-OCT-2003
                                               ABM53777;
                                                                                     ABM53777
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밁 á

standard; protein; 112

0

(first entry)

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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                      nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infections. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (BLISA). Note: The sequence data for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis uveitis; endophthalmitis; bone; joint; central nervous system; ELISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-APR-2000; 2000US-0199047P
02-JUN-2000; 2000US-0208841P
07-JUL-2000; 2000US-0216747P
                                                                                                                                                                                                                                                             Sequence 112 AA;
                                                                                                                                                                                                                                                                                                                        this patent did not form part of the printed specification, obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and osteomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO 18453; 1069pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L'maisonneuve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-APR-2001; 2001WO-US012865
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CORI-) CORIXA CORP.
                                                                                                                             Local Similarity
ses 6; Conserv
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35
                                                           1 FSRYAR
FSRYAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             acne vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Persing DH, I
                                                                                                                                Conservative
   40
                                                                                                                                                            100.0%;
                                                                                                                                                            100.0%; Score 31; 100.0%; Pred. No.
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, Jen S, Carter D;
                                                                                                                                0,
                                                                                                                                   Mismatches
                                                                                                                                                               39,
                                                                                                                                                                                               DB 4;
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al nervous system; ELISA;
                                                                                                                                   0
                                                                                                                                                                                               Length 112;
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                                                                                                                                                                                                                      CC polynuclectide of the invention; antibodies against polyneptides of the invention; fusion proteins comprising a polypeptide of the invention; a comment of the invention of an immune response specific for a p. acrose composition of an immune response of the invention of the invention. The polyment of an immune response of the invention of the invention of the invention of the invention. The polyment of an immune response of the invention of the invention of the invention. The polyment of an immune response of the invention of an immune response of the invention. The polyment of the vaccine composition is useful for the contained within the p. acros, or for treating acros, and the kit is useful for performing a diagnostic assay. The present contained within the p. acros polyment of the present of the present of the present of the present of the printed specification, but was obtained in electronic format directly from wipo at the present of the present o
                                                                                             Matches
                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mitcham
Zhang Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  encoding a Propionibacterium acnes procedure. AND 18864536) and to polypeptides encoded by the polypucleotides (ABM35624-ABM64536) and to immunogenic fragments of P. acnes polypeptides. The invention additionally encompasses expression vectors and host cells comprising a additionally encompasses expression vectors and host cells comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                        Sequence 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-OCT-2001; 2001US-00978825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acne vulgaris; antiseborrhoeic; dermatological; antibacterial; immunostimulant; immune response; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium acnes predicted ORF-encoded polypeptide #18453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CORI-)
                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to an isolated polynucleotide (ACF64435-ACF64733) oding a Propionibacterium acnes protein. The invention also relates to
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                                                                                           Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; SEQ ID NO 18453; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACF64510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORIXA
FSRYAR
                                            FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                L, Skeiky YAW, Pe
Wang S, Jen S, I
Vallieve-Douglass
                                                                                             Conservative
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                                                                                                                 100.0%;
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Lodes MJ,
                                                                                           0;
                                                                                        Score 31; DB
Pred. No. 39;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bhatia A,
Benson DR,
                                                                                                                                         6
                                                                                           0;
                                                                                                                                       Length 112;
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Jones R, Carter
                                                                                        0,
                                                                                           Gaps
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RESULT 14 ABM64633 ID ABM64

ABM64633 standard; protein;

174 AA

밁 8

35 FSRYAR

Query Match Best Local S Matches 6

Similarity 6; Conser FSRYAR 6

Conservative

0

Mismatches

6 0

Length 174; Indels

0,

Gaps

0

100.0%;

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CC encoding a Propionibacterium acres protein. The invention also relates to cold polypeptides encoded by the polypucleotides (ABM35624-ABM64536) and to CC immunogenic fragments of P. acres polypeptides. The invention also relates to CC immunogenic fragments of P. acres polypeptides. The invention of comprising a polypeptide of the invention of the invention; antibodies against polypeptides of the invention; antibodies against polypeptides of the invention; a comprising a polypeptide of the invention; a comprising a polypeptide of the invention; a cc polypeptide and an isolated T cell population comprising T cells prepared complete the comprising P. acres polypeptides, antigen-presenting cells that express the polypeptide; or comprising P. acres in a cc patient; and a method for inhibiting the development of P. acres in a cc patient; and a method for inhibiting the development of P. acres in a cc patient. The P. acres polypeptides, polypucleotides, antibodies, fusion proteins. T cell populations or antigen-presenting cells that express the collypeptides are useful for diagnosing, preventing or treating acre composition. The polymucleotides can also be used as probes or primers for comprising or for a P. acres for complete acid hybridisation. The vaccine composition is useful for the contain an immune response specific for a P. acres conditions or antigen-presenting cells that express the composition of an immune response against P. acres, or for treating acre, can be contained present as probes or primers for composition of an immune response against P. acres, or for treating acre, this patent did not form part of the printed specification, but was contained present of the printed specification.
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Barth B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris, or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                   Sequence 174 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an isolated polynucleotide (ACF64435-ACF64733)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 7; SEQ ID NO 29309; 1481pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Propionibacterium acnes immunogenic polypeptide #29309
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                                                                                                                                                                                                                     .int/pub/published_pct_sequences
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Lodes MJ,
Score 31; DB
Pred. No. 61;
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Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maisonneuve JL;
Jones R, Carter
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ABB90579
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Best Local Similarity
                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see ANY34584-Y35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae. (Updated on 17 -OCT-2003 to standardise OS field)
Chlamydial infection; antigen; immunogen; vaccine; diagnosis; human respiratory disease; cardiovascular disease; atherosclerosis; coronary artery disease; carotid artery stenosis; myocardial infarction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine;
                                                                                               29-AUG-2003
29-JUL-2002
                                                                                                                                                                  ABB90579 standard; protein; 357
                                                                                                                                                                                                                                                                                                                                                                      Sequence 357 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY34584-Y35879 represent the proteins encoded by all the open reading frames in the complete genome (see AAX91990) of Chlamydia pneumoniae. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-NOV-1998;
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13-SEP-1999
                                                              Chlamydia pneumoniae cp7224 protein, SEQ ID NO:107
                                                                                                                                          ABB90579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 1303-1304; Disclosure; 1912pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-357842/30.
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                                                                                                                                                                                                                                                135 FSRYAR 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence of Chlamydia pneumoniae.
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                                                                                                                                                                                                                                                                                                           Conservative
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(first en
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                                                                                              (revised)
(first entry)
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98US-0107078P
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                                                                                                                                                                                                                                                                                                        100.0%; Score 31; DB 2; 1 100.0%; Pred. No. 1.3e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                       Length 357;
                                                                                                                                                                                                                                                                                                           Indels
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RESULT 17
ABU27032
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                                                                                                                                                                                                                                                                                                                                 CC Sequences ABB90526-ABB90715 represent novel proteins from Chlamydia CC pneumoniae (strain CWLO29), and ABL91184-ABL91373 represent DNA encoding Ct them. The proteins are predicted to be immunogenic and may therefore be useful in vaccine production and for diagnostic purposes. Chlamydia CC pneumoniae is a common cause of respiratory disease in humans, and is CC also involved in the development of cardiovascular diseases such as CC atherosclerosis, coronary artery disease, carotid artery stenosis, CC myocardial infarction, cerebrovascular disease, aortic aneurysm, CC claudication and stroke. The proteins and nucleic acids of the invention CC may be used in vaccines and pharmaceutical compositions for the prevention or treatment of chlamydial infections, particularly Chlamydia pneumoniae infections. The proteins may also be used in the detection of CC chlamydia pneumoniae, and the nucleic acids may be used in pCR, branched CC pneumoniae gene expression. The present sequence represents a specifically claimed Chlamydia pneumoniae protein of the invention. CC (Updated on 29-AUG-2003 to standardise OS field)
                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                             Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JUL-2000;
11-JUL-2000;
21-JUL-2000;
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14-SEP-2000;
                                                                                                                                                                                                                                                                                                          Sequence 357 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel Chlamydia pneumoniae protein useful in the manufacture of a medicament for treatment or prevention of infection due to Chlamyc preferably Chlamydia pneumoniae, and for diagnostic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cerebrovascular disease; aortic aneurysm; claudication; stroke;
strain CWL029.
            Protein encoded by Prokaryotic
                                         23-OCT-2003
19-JUN-2003
                                                                                      ABU27032
                                                                                                                ABU27032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 93; 364pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-2000;
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                                                                                                                                                                                                                                              Similarity 6; Conserv
                                                                                                                                                                                                            FSRYAR 6
                                                                                                                  standard;
                                                                                                                                                                                        FSRYAR 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ); 2000GB-00016363.
; 2000GB-00017947.
; 2000GB-00017963.
; 2000GB-00019368.
; 2000GB-00020440.
; 2000GB-00022583.
; 2000GB-00027549.
; 2000GB-00031706.
                                                                                                                                                                                                                                                 Conservative
                                         (revised)
(first entry)
                                                                                                                protein;
                                                                                                                                                                                                                                                               100.0%;
                                                                                                                  357
                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                             Score 31; DB 5;
Pred. No. 1.3e+02;
             essential
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                                                                                                                                                                                                                                                                           Length 357;
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gene

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                                                                                                                                                                                                                                                                                                                                                                                                 cc encoding a polypeptide whose expression is inhibited by the antisense compoling a polypeptide whose expression is inhibited by the antisense cc polypeptide or its fragment whose expression is inhibited by the cartier acid; (4) an antibody capable of specifically binding ct the polypeptide; (5) producing the polypeptide; (6) inhibiting cellular cc proliferation or the activity of a gene in an operon required for crown product or that has an activity against a biological pathway crequired for proliferation, or that inhibits cellular proliferation; (8) clentifying a compound that inhibits proliferation; (8) clentifying a gene required for cellular proliferation or the biological pathway crequired for proliferation, or that inhibits cellular proliferation of the biological pathway in which a proliferation-required gene or its gene product lies or a gene on which the test compound that inhibits proliferation of an organism acts; (9) manufacturing an antibiotic; (10) profiling a compound's activity; (11) a culture comprising strains in which the gene compound's activity; (11) a culture comprising strains in which the gene compound's activity; (11) a culture comprising strains in which the gene compound's activity; (11) a culture comprising strains in which the gene compound that inhibits the compound that inhibits the compound that inhibits the condition of an organism. The antisense nucleic acids are useful for cidentifying or screaming for homology.
                                                                                Matches
                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wang
Wall
                                                                                                                                                                                                                                            identifying proteins or screening for homologous nucleic acids required for cellular proliferation to isolate candidate molecules for rational drug discovery programs, or for screening homologous nucleic acids required for proliferation in cells other than S. aureus, S. typhimurium, K. pneumoniae or P. aeruginosa. The present sequence is encoded by one of the target prokaryotic essential genes. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-OCT-2001;
08-FEB-2002;
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06-SEP-2001;
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                                                                                                                                                             Sequence
                                                                                                                                                                                                        ftp.wipo.int/pub/published_pct_sequences. (Updated on 23-OCT-2003 to
standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense nucleic acids, useful for identifying proteins or screening for homologous nucleic acids required for cellular proliferation to isolate candidate molecules for rational drug discovery programs.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                e 6213 antisense sequences given in the specification where expression the nucleic acid inhibits proliferation of a cell. Also included are:

) a vector comprising a promoter operably linked to the nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to an isolated nucleic acid comprising any one of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ם בֿי
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                                                                                                    Similarity
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Trawick
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                                          FSRYAR 6
FSRYAR 140
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                                                                                Conservative
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2002US-00072851.
2002US-0362699P.
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2001US-00948993.
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Carr GJ,
                                                                                                  100.0%; Score 31; DB 6; 100.0%; Pred. No. 1.3e+02;
                                                                                0
                                                                              Mismatches
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Forsyth
                                                                                                                     Length 357;
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Xu HH;
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RESULT 19 ADS44828

ADS44828

ADS44828 standard; protein; 463

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RESULT 18
AEC95703
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                                                                                                                                                                                                                                                                                                                                                         The invention relates to a polypeptide for use as an autotransporter cantigen. Also provided is a method of eliciting an immune response in an individual which involves administering to the individual an Chlamydia pneumoniae autotransporter protein selected from Cpn0794, Cpn0795.

Cpn0796, Cpn0797, Cpn0798, or Cpn0799 or an immunogenic fragment. The polypeptide is secreted into the cytoplasm of the host cell through a Crype v autotransporter secretion system mechanism. The polypeptide is useful as an antigen for raising a C. pneumoniae specific immune response in an cresponse. It is also useful for raising a systemic immune response in an individual infected with C. pneumoniae. The polypeptide is diagnosis. It can also be used in the preparation of a medicament for the prevention or treatment of a C. pneumoniae infection in an individual. It can also be used as an autotransporter protein which immunoreacts with seropositive serum of an individual infected with C. pneumoniae. The preparation of an individual in the preparation of an individual in the preparation of an individual. The preparation of an autotransporter protein which C. pneumoniae. The preparation fan individual infected with Seropositive serum of an individual infected with C. pneumoniae. The preparation (pn anda)
                                                                                                                                  Matches
                                                                                                                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                    Sequence 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Chlamydia pneumoniae autotransporter protein, useful as an autotransporter antigen, for raising a specific immune responso preventing or treating C. pneumoniae infections in individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-MAR-2004; 2004US-0549832P.
12-JAN-2005; 2005US-0643110P.
19-JAN-2005; 2005US-0644552P.
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                                                                                                                                                                                                                                                                                                                                    pneumoniae MurG peptidoglycan transferase protein (CPn 0904).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENBANK;
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135 FSRYAR 140
                                                               1 FSRYAR 6
                                                                                                                                  Similarity 6; Conserv
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                                                                                                                                     Conservative
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                                                                                                                                                              100.0%;
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                                                                                                                                  Score 31; DB 9; Pred. No. 1.3e+02; Mismatches 0;
                                                                                                                                                                                                 Length 357;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immune response
                                                                                                                                         Indels
                                                                                                                                     0
                                                                                                                                  Gaps
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02-DEC-2004

(first entry)

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                                                                                                                                                                     CC having an improved property comprises transforming a plant with the CC recombinant DNA construct and growing the transformed plant, where the CC polynucleotide or polypeptide is useful for improving plant properties. CC The recombinant DNA construct is useful for producing plant properties. CC improved plant properties, e.g. improved cold, heat or drought tolerance, tolerance to herbicides, extreme osmotic conditions, pathogens or pests, cc increased resistance to plant disease, better growth rate by modification of the cell cycle pathway with plant growth regulators, increased rate of CC content, improved yield by modification of carbohydrate, nitrogen or CC content, improved plant growth and development under at least one stress crowding improved plant growth and development under at least one stress CC condition, improved lignin production of photosynthesis or by CC providing improved plant growth and development under at least one stress CC condition. This sequence represents a bacterial polypeptide used in the scope of the invention. Note: The sequence data for this patent did not CC form part of the printed specification but was obtained in electronic format from USPTO at sequence.
                                                                                        Query Match
Best Local :
                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cold tolerance; heat tolerance; drought tolerance; herbicide; osmosis; pathogen tolerance; pest tolerance; plant disease resistance; cell cycle pathway modification; plant growth regulator; homologous recombination; seed oil yield; protein yield; carbohydrate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                microbial source. The invention also relates to a transformed plant comprising the recombinant DNA construct and a method of producing a transformed plant having an improved property. The plant is a crop plant such as maize or soybean. The method of producing a transformed plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a recombinant DNA construct comprising a promoter functional in a plant cell, where the promoter is positioned to provide for expression of a polynucleotide encoding a polypeptide from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New recombinant DNA construct comprising a promoter positioned to provi
for expression of a polynucleotide encoding a polypeptide from a
microbial source, useful for producing plants with improved properties.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacterial polypeptide #23258.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 23258; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-061375/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HINK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-FEB-2002; 2002US-0360039P
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                                                                                                                                            Sequence 463
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                                                                                                                                                          congruent to secondary structural elements in the other family, so as to determine if the families are related by common ancestry or not. The method is used to confirm/deny the hypothesis that proteins are homologous and related methods are used to identify mutations during divergent evolution of proteins, to identify in vitro properties of proteins that are important for physiological activity and to generate genome-sized databases. The present sequence is Escherichia coli 6-phospho-strand-glucosidase (EC 3.2.1.86). This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                             The invention relates to a method for excluding homology between two protein families. The method involves constructing models for secondary structural elements for each family; aligning secondary structural elements from the elements of one family with the secondary structural elements from the other family around sequence motifs; determining whether secondary structural elements flanking the sequence motifs in one family are
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Methods for excluding or detecting homology between protein families, useful e.g. for identifying in vitro properties of proteins important physiological activity.
                                                                                                                                Sequence 466 AA;
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AAU34677 standard; protein; 474

AAU34677;

14-FEB-2002 (first entry)

coli cellular proliferation protein #258

Antisense; prokaryotic cellular proliferation protein; antibiotic; antibacterial; drug design.

coli.

WO200170955-A2

27-SEP-2001

21-MAR-2001; 2001WO-US009180

RESULT 21
AAU34677
ID AAU34
XX 21-MAR-2000; 23-MAY-2000; 26-MAY-2000; 23-OCT-2000; 27-NOV-2000; 22-DEC-2000; 2000US-0191078P. 2000US-0206848P. 2000US-0207727P. 2000US-0242578P. 2000US-0253625P. 2000US-02537931P.

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RESULT 22
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                                                                                                                                                                                                                                                                                                                                                                                                     pathogen tolerance; pest tolerance; plant disease resistance;
cell cycle pathway modification; plant growth regulator;
homologous recombination; seed oil yield; protein yield; carbohydrate;
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                                21-FEB-2002; 2002US-0360039F
                                                                                           20-FEB-2003; 2003US-00369493.
                                                                                                                                                         18-DEC-2003
                                                                                                                                                                                                                       US2003233675-A1
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les 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 FSRYAR 6
                                                                                                                                                                                                                                                                                                                                             polypeptide
                                                                                                                                                                                                                                                                                                                                                                    phosphorus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 31; DB 4; I ilarity 100.0%; Pred. No. 1.7e+02; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A,
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Xu HH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ŗ
                                                                                                                                                                                                                                                                                                                                                                       photosynthesis; lignin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wall D,
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ucleic acids
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18-FEB-1998; 18-FEB-1999;

98US-0074788P 99US-00252991 US6551795-B1

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RESULT 23
ABO74595
ID ABO74
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AC ABO74
XX
DT 29-JU
DX
DX
Pseud
XX
Bacte
XX
Bacte
XX
US655
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PS 18-FI
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18-FI
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Best Local Similarity
Watches 6; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising the recombinant DNA construct and a method of producing a crop plant of transformed plant having an improved property. The plant is a crop plant country as maize or soybean. The method of producing a transformed plant construct and as maize or soybean. The method of producing a transformed plant construct and growing the transformed plant with the crop plant comprises transformed plant with the crop plant construct and growing the transformed plant with the construct and growing the transformed plant where the conjunction of the construct is useful for improving plant properties. The recombinant DNA construct is useful for producing plant properties. Increased resistance to harbicides, e.g. improved cold, heat or drought tolerance, increased resistance to plant disease, better growth rate by modification of the cell cycle pathway with plant growth regulators, increased rate of the monologous recombination, modification of carbohydrate, nitrogen or phosphorus use and/or uptake, by modification of photosynthesis or by growiding improved plant growth and development under at least one stress condition, improved plant growth and development under at least one stress condition, improved plant growth as bacterial polypeptide used in the groups of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic form part for the granted specification but was obtained in electronic condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New recombinant DNA construct comprising a promoter positioned to provi
for expression of a polynuclectide encoding a polypeptide from a
microbial source, useful for producing plants with improved properties.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CAOY/)
(HINK/)
(SLAT/)
(CHEN/)
(GOLD/)
                                                                                                                                                                                                                                                                                    ABO74595;
                                                                                                                                                                                                                                                                                                                     ABO74595 standard; protein; 509 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 474 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      promoter functional in a plant cell, where the promoter is positioned in a polynucleotide encoding a polynucleotide from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-061375/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cao Y,
                                                                                                                                                                               Bacterial infection;
                                                                                                                                                                                                             Pseudomonas aeruginosa polypeptide #6770.
                                                                                                                                                                                                                                                  29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a recombinant DNA construct comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 23518; 122pp; English.
                                                                                                                                           Pseudomonas aeruginosa
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SLATER S
CHEN X.
GOLDMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                     1 FSRYAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              from USPTO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                               Pseudomonas aeruginosa infection; antibacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 31; 100.0%; Pred. No. 1
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1.7e+02;
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RESULT 24
ANBC24444
ID ANBC24
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AC ANCA

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs, including anti-P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa-derived peptides or polypeptides, as target components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypeptides of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at segdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
N-PSDB;
                                                                                      Drmanac
                                                                                                                                                                                                  31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                     30-MAR-2001; 2001WO-US008631.
                                                                                                                                                                                                                                                                                                                                                 11-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABG24444;
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                                                                                                                                         (HYSE-) HYSEQ INC.
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                             2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  n; chromosome mapping; gene mapping; gene therapy; forensic;
supplement; medical imaging; diagnostic; genetic disorder.
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                                                                                   Liu C,
                                                                                                                                                                                                  2000US-00540217.
2000US-00649167.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                   Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein #24435.
                                                                                         ΥŢ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 31; DB 7;
Pred. No. 1.8e+02;
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Length 1225;

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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain (C reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used (C in diagnostics as expressed sequence tags for identifying expressed (C genes. (I) is useful in gene therapy techniques to restore normal (C activity of (II) or to treat disease states involving (II). (II) is (C useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food (C supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders (involving aberrant protein expression or biological activity. The (C involving aberrant protein expression or biological activity. The (C diagnostics, forensics, gene mapping, identification of mutations (C amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in (C entertonic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 20; SEQ ID NO 54803; 103pp; English
Sequence 1225 AA;
                                              ftp.wipo.int/pub/published
                                              _pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seful in mutations
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RESULT 25 ABG24819 ID ABG24 Matches Query Match Best Local 31-MAR-2000; 23-AUG-2000; 30-MAR-2001; 2001WO-US008631 Human; chromosome mapping; Novel human diagnostic protein #24810 ABG24819 standard; protein; 1258 AA. WO200175067-A2 Homo sapiens. 18-FEB-2002 supplement; 613 FSRYAR 618 1 FSRYAR 6 Similarity 6; Conserv RT, HYSEQ Liu C, Conservative 2000US-00649167 2000US-00540217. (first entry) INC. medical 100.0%; g; gene mapping; gene therapy; forensic imaging; diagnostic; genetic disorder. ΥŢ 0 Score 31; DB 4; Pred. No. 4.5e+02; Mismatches 0, Indels 0, Gaps

N-PSDB; AAS89006. WPI; 2001-639362/73

New isolated polynucleotide and

encoded polypeptides,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                           SAPHO syndrome; synovitis; acne; pustulosis; hypertosis; osteomyelitis; uveitis; endophthalmitis; bone; joint; central nervous system; EIISA; inflammatory lesion; acne vulgaris; enzyme linked immunosorbent assay; dermatological; osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymerlection are also used
WPI; 2001-616774/71.
N-PSDB; AAS59612.
                                                                                                                                           21-APR-2000; 2000US-0199047P.
02-JUN-2000; 2000US-0208841P.
07-JUL-2000; 2000US-0216747P.
                                                                                                                                                                                                                                                                                         WO200181581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Propionibacterium acnes immunogenic protein #21584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU60688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU60688 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1258 AA;
                                                        L'maisonneuve
                                                                                                                                                                                                               20-APR-2001; 2001WO-US012865.
                                                                                                                                                                                                                                                     01-NOV-2001
                                                                                                                                                                                                                                                                                                                           Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in diagnostics as expressed sequence tags for identifying expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention relates to isolated polynucleotide (I) and polypeptide (II) usinces. (I) is useful as hybridisation probes, polymerase chain
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                                                                                                           CORIXA CORP
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                                                                       Persing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO 55178; 103pp; English
                                                                                                                                                                                                                                                                                                                             acnes.
                                                      Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                       PH,
                                                        ĸ.
                                                        Mitcham JL, Wang SS,
, Jen S, Carter D;
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Pred. No. 4.6e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1258;
                                                                         Bhatia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to downregulate expression and activity of P. acnes polypeptides and therefore treat P. acnes infectiois. The antibodies may also be used as diagnostic agents for determining P. acnes presence, for example, by enzyme linked immunosorbent assay (BLISA). Note: The sequence data for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful for treating acne vulgaris.
                                                                                               Mitcham C
Zhang Y,
Barth B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 58 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the treatment, prevention and diagnosis of medical conditions carre, P. acnes. The disorders include SAPHO syndrome (synovitis, acne, p. acnes, hypertosis and contecomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central p. acnes is also involved in infections of bone, joints and the central p. acnes is also involved in infections of bone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAU39105-AAU68017 represent Propionibacterium acnes immunogenic polypeptides. The proteins and their associated DNA sequences are used in the treatment, prevention and diagnosis of medical conditions caused by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM57207;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM57207 standard; protein; 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           this patent did not form part of the printed specification, obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ ID NO 21883; 1069pp; English
                                                WPI; 2003-381789/36
N-PSDB; ACF64541.
                                                                                                                                                                                                                                                                                                                                         Propionibacterium acnes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-OCT-2003
                                                                                                                                                                                                  15-OCT-2001; 2001US-00978825
                                                                                                                                                                                                                                    11-OCT-2002; 2002WO-US032727
                                                                                                                                                                                                                                                                        24-APR-2003
                                                                                                                                                                                                                                                                                                       WO2003033515-A1
                                                                                                                                                                                                                                                                                                                                                                                          Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                          Propionibacterium acnes predicted ORF-encoded polypeptide #21883.
                                                                                                                                                                   (CORI-)
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5; Conserv
                                                                                               JL, Skeiky YAW, Pers
, Wang S, Jen S, Lod
, Vallieve-Douglass J;
                                                                                                                                                                   CORIXA CORP
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                                                                                                                                                                                                                                                                                                                                                                           response; vaccine.
                                                                                                                   Persing DH,
Lodes MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 28; DB
Pred. No. 95;
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                                                                                                                   Bhatia
Benson [
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                                                                                                                     DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 58;
                                                                                                                 Maisonneuve JL;
Jones R, Carter
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New Propionibacterium acnes polypeptides and polynucleotides encoding the polypeptide, useful for diagnosing, preventing or treating acne vulgaris,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                 Human; secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping; signal peptide; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopoiesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition.
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  Dumas Milne Edwards
                                                                               01-AUG-1997;
                                                                                                                     31-JUL-1998;
                                                                                                                                                                                                      WO9906554-A2
                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY12203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY12203 standard; protein; 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 58 AA;
                                      (GEST ) GENSET
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5; Conserv
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                                                                               97US-00905134
                                                                                                                       98WO-IB001238.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  protein
Duclert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 28; DB Pred. No. 95; 1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID
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Lacroix
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RESULT 29
ABR40679
ID ABR40
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DRX PP11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                       Allen SM,
Jones TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acids encoding human secreted proteins - obtained from cDNA libraries prepared from kidney, fetal kidney, dystrophic muscle, muscle and heart tissue.
                                                                                                                                                                                                                                                                                                       Plant; oil trait; oil phenotype; altered lipid profile; MAP kinase receptor-like protein kinase; mitogen activated protein kinase; oi LIP15-like transcription factor caleosin; ATP citrate lyase; SNF1;
                                                                                                                                                                                                                                                Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 85
                                                                                    (DUPO )
                                                                                                                                                          27-JUN-2002; 2002WO-US020152
                                                                                                                                                                                         09-JAN-2003
                                                                                                                                                                                                                                                                             transgenic plant.
                                                                                                                                                                                                                                                                                         CKC-like transcription factor; antisense inhibition; co-suppression;
                                                                                                                                                                                                                                                                                                                                                                                                  16-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR40679 standard;
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                           Tarczynski
                                                                                                                             29-JUN-2001; 2001US-0301913P
                                                                                                                                                                                                                     WO2003002751-A2
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                                      Allen V
Kinney
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                                                                                                                                                                                                                                                                                                                                                                   oil trait related protein sequence SEQ
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                                       Υ ΑJ,
                                                                                                  NEMOURS &
                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein; 90
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                                         Cahoon
Klein
                                                                                    INT INC
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                                         J R
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Pred. No. 1.4e+02;
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                                       Epelbaum S, Fan
Li C, Oliveira
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                                         Famodu OO, H
ira IC, Sakai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 85;
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                                                                                                                                                                                                                                                                                                                                                                     ID NO:178
                                                                                                                                                                                                                                                                                                                                       MAP kinase;
                                                        Harvell LT;
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WPI; 2003-201509/19

N-PSDB; ACC00714.

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ARESULT 30
ABR40677
ID ABR40677
AC ABR40
XX ABR40
AC ABR4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                           (DUPO )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plant; oil trait; oil phenotype; altered lipid profile; MAP kinase; receptor-like protein kinase; mitogen activated protein kinase; oil; LIP15-like transcription factor caleosin; ATP citrate lyase; SNF1; CKC-like transcription factor; antisense inhibition; co-suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oryza sativa oil trait related protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 12; Page 215;
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                                                                                                                                           Allen SM, Allen WB,
Jones TJ, Kinney AJ,
Tarczynski MC;
                                                                                                                                                                                                                                                                                                                                                                                                 29-JUN-2001; 2001US-0301913P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-JUN-2002; 2002WO-US020152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transgenic plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR40677 standard; protein; 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phenotypes in plants such as sunflower,
                                         2003-201509/19
DB; ACC00712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
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                                                                                                                                                                                                                                                                                           DU PONT DE NEMOURS & CO
PIONEER HI-BRED INT INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 83.: 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FSKYAR 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        542pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.3%;
                                                                                                                                                                                 Cahoon
Klein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 28; DB 6; Le
Pred. No. 1.5e+02;
1; Mismatches 0;
                                                                                                                                                                                 J RE
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                                                                                                                                                                                 Epelbaum S, Fan
Li C, Oliveira
                                                                                                                                                                                                                                                                                                                                 H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 90;
                                                                                                                                                                             Famodu OO, Harvell LT;
ira IC, Sakai H, Shen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NO:174.
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RESULT 31
AAW67901
ID AAW67901
XX AAW67
XX AAW67
XX Human
XX Human
XX Human
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Cogni
XW inmur
XW cogni
XW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes an isolated nucleotide fragment (I) comprising a nucleic acid sequence (NS) chosen from a NS encoding a polypeptide (PP) having receptor-like protein kinase activity, mitogen activated protein (MAP)-kinase activity, LIPIS-like transcription factor activity and CKC-like transcription factor activity and CKC-like transcription factor activity. Also described: (I) complement (II) of (I); (2) a chimeric construct (III) comprising (I) or (II), operably linked to a regulatory sequence; (3) a plant (IV) comprising (III) in its genome; (4) seeds (V) obtained from (IV); and (5) constrained from (V). (I) or its part can be used in antisense inhibition or co-suppression in a transformed plant. (III) is useful for canola, Brassica, sorghum, sunflower or coconut. (III) is also useful for creating transgenic plants having altered lipid profiles. (I) can also be used as a hybridisation probe. ACC00626 to ACC00868 and ABR40591 to ABR40879 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 5
21-MAR-1997;
21-MAR-1997;
21-MAR-1997;
21-MAR-1997;
30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel nucleotide fragment encoding polypeptides having receptor-like protein kinase activity, caleosin-like activity, useful for altering phenotypes in plants such as sunflower, coconut, soybean, wheat and 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW67901 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 132 AA;
                                                                                                                                                                                                                                                19-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted protein encoded by gene 36 clone HODCL36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-1999
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                         97US-0041276P.
97US-0041277P.
97US-0041281P.
97US-0042344P.
97US-0048069P.
97US-0048094P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                /label= unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 542pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.3%;
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6; I
2.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    soybean, wheat and rice
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Best Local Similarity
Thes 5; Conserve
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30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                   This sequence represents a secreted human protein encoded by the gene clone detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin FC portion (e.g. AAX00602) for increasing the stability of the fused protein as compared to the human protein only. The invention relates to 87 novel genes and their fragments (nucleic acid sequences: AAX00611-X00724; amino acid sequences AAW67807-W68004) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypucleotides in a sample or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the 87 polynucleotides, based on which tissues they are most highly expressed in (see AAX00611 for described uses)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Young P, Florence
WO2003048304-A2
                        Alloiococcus otitis
                                                              Alloiococcus
                                                                                                                  20-NOV-2003
                                                                                                                                                                    ADB06538 standard; protein; 145 AA
                                                                                                                                                                                                                                                                                                                                               Sequence 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorders, immune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-070066/06.
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19-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI
                                               iococcus otitidis; antigenic protein; immunogenic;
therapy; Gram-positive bacterium; infection.
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                                                                                                                                                                                                                                                                 FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Greene JM,
KA, Olsen I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 329;
                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                               A,
                                                                                     otitis antigenic protein
                                                                                                                 (first entry)
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97US-0048188P.
97US-0048351P.
97US-0048352P.
97US-0048355P.
97US-005037P.
97US-0054804P.
97US-0056370P.
97US-0056862P.
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97US-0048099P.
97US-0048131P.
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97US-0048154P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             385pp; English.
                                                                                                                                                                                                                                                                                                     90.3%;
83.3%;
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S, Ebner R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    eatment of e.g. cancers, neurological inflammation or blood disorders.
                                                                                                                                                                                                                                                                                         Score 28; DB 2;
Pred. No. 2.3e+02;
1; Mismatches 0
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Brewer LA,
                                                                                        SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosen CA, bu
                                                                                        NO:478
                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                   Length 139
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A, Shi Y;
                                                              immunisation;
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                                                                                                                                                                                                                                                                                          Gaps
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RESULT 33 ADB06540

ADB06540 standard; protein; 172

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SXXXXXXXXXXXXX

Alloiococcus

Alloiococcus otitidis; antigenic protein; immunogenic; immunisation; gene therapy; Gram-positive bacterium; infection.

otitis antigenic protein SEQ ID NO:480.

(first entry)

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                                                                                                                                              CC expression vector comprising the novel isolated polynucleotide (1), its complement, degenerate variant or fragment; (3) a genetically engineered (2) host cell, transfected, transformed or infected with the vector of (2); (CC (4) an antibody specific for the polypeptide of (1); (5) an immunogenic (CC composition comprising the polypeptide of (1); (5) an immunogenic (CC composition comprising the polypeptide of (1); (5) an immunogenic (CC polypeptide of (1) and a carrier; (7) a protein chip comprising the (CC polypeptide of (1) and a carrier; (7) a protein chip comprising an array (CC of the polypeptides of (1), their biological equivalent or fragment; (8) (CC immunising against Alloicoccus otitidis by administering to a host the continuing the novel polypucleotide, its degenerate variant or fragment; (CC or the antibody of (4); and (11) producing a polypeptide by culturing the (CC polypeptide from the culture. (1) can be used in gene therapy. The (CC polypeptides, polypeptides, antibodies and compositions of the present invention can be used for treating and diagnosing diseases, drug (CC stridies). The present sequence represents an Alloicoccus otitidis. The present sequence represents an Alloicoccus otitidis.
                                                                      Matches
                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Alloiococcus otitidis polynucleotides and polypeptides, treating and diagnosing diseases, drug screening assays and effects during drug clinical trials.
                                                                                                                                            Sequence 145 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an isolated polypeptide that is encoded by the polynucleotide (I); (2) an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 33; SEQ ID NO 478; 1019pp; English.
                                                                                                                                                                                antigen protein from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alloiococcus otitidis genomic DNA, which encodes an antigenic protein Alloiococcus otitidis is a Gram-positive bacterium. Also described: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes an isolated polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fletcher LD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-NOV-2001;
18-NOV-2002;
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122 FSRYSR 127
                                    1 FSRYAR 6
                                                                    Similarity
5; Conserv
                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2001US-0333777P.
2002US-0426742P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HOLDINGS CORP
                                                                                       90.3%;
                                                                    Score 28; DB 6;
Pred. No. 2.4e+02;
1; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Russell DP,
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                                                                                                        Length 145;
                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , useful for d monitoring
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of.

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RESULT 34
AAW67842
ID AAW6
XX
AC AAW6
XX
DT 25-M
XX
DE Huma
XX
KW Huma
KW diag
                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an isolated polynucleotide (I) of CC Alloiococcus otitidis genomic DNA, which encodes an antigenic protein. CC Alloiococcus otitidis is a Gram-positive bacterium. Also described: (1) CC an isolated polypeptide that is encoded by the polynucleotide (I); (2) an CC expression vector comprising the novel isolated polynucleotide (I); (2) an CC complement, degenerate variant or fragment; (3) a genetically engineered CC composition comprising the polypeptide of (1); (5) an immunogenic composition comprising the polypeptide that is comprised in the CC expression vector; (6) a pharmaceutical composition comprising the polypeptide that is comprised in the CC polypeptide of (1) and a carrier; (7) a protein chip comprising an array of the polypeptides of (1), their biological equivalent or fragment; (8) immunising against Alloiococcus otitidis by administering to a host the continist in the biological sample; (10) a kit comprising a container CC containing the novel polynucleotide, its degenerate variant or fragment; or the antibody of (4); and (11) producing a polypeptide by culturing the CC polypeptide from the culture. (I) can be used in gene therapy. The CC polynucleotides, polypeptides, antibodies and compositions of the present invention can be used for treating and diagnosing diseases, drug creening assays and monitoring of effects during drug clinical trials. The present sequence represents an Alloiococcus otitidis
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                                                                                                                                                                                                                                                문
                                                                                                                                                                                                                                                                                                                      Matches
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18-NOV-2002; 2002US-0426742P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 33; SEQ ID NO 480; 1019pp; English.
Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia;
                                                                                                                                                           AAW67842 standard; protein; 174 AA
                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                     Human secreted
                                                                                                                          AAW67842
                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                      (first entry)
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                                                     protein
                                                                                                                                                                                                                                                                                                                                    90.3%;
                                                   encoded by gene 36 clone HODCL36.
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                                                                                                                                                                                                                                                                                                                    Score 28; DB 6;
Pred. No. 2.9e+02;
1; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zagursky
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                                                                                                                                                                                                                                                                                                                                                     Length 172
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30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
30-MAY-1997;
05-AUG-1997;
19-AUG-1997;
02-OCT-1997;
                        This sequence represents a secreted human protein encoded by the gene clone detailed in the descriptor line. The gene can be used to generate fusion proteins by linking to the gene to a human immunoglobulin FC portion (e.g. AAX00602) for increasing the stability of the fused protein as compared to the human protein only. The invention relates to 87 novel genes and their fragments (nucleic acid sequences: AAX00611-X00724; amino acid sequences AAAW67807-M68004) which are useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. Also, pathological conditions can be diagnosed by determining the amount of the new polypucleotides in a sample or by determining the presence of mutations in the new polypucleotides. Specific uses are described for each of the 87 polypucleotides, based on which tissues they are most highly expressed in a sample or an expression of the second control of the second contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 developmental abnormality; foetal deficiency; blood; allergy; renal; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obseity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders.
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30-MAY-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 11; Page 291; 385pp; English.
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polynucleotides, (see AAX00611 for
        AAX00611 for
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97US-0048187P.
97US-0048188P.
97US-0048351P.
97US-0048352P.
97US-0048355P.
97US-0050937P.
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97US-0048095P.
97US-0048096P.
97US-0048099P.
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97US-0048154P.
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            described uses)
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S, Ebner R,
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Brewer LA,
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Shi Y;
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Query Match Best Local S Matches 5

Similarity 83.: 5; Conservative

90.3%;

Score 28; DB 2; Lo Pred. No. 2.9e+02; 1; Mismatches 0;

Length 174;

Indels

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Sequence

AA;

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RESULT 36
AEP69875
ID AEP69
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XX Micrc
XX Satel
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                                                                                                                                                                                                               Matches
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Best Local (
                                                                                                                                                                                                                                                                                                The invention relates to an isolated nucleic acid encoding a PR polypeptide. The polypeptide, agonist or an antagonist, antibod composition, and method are useful for diagnosing and treating related disorder, e.g. systemic lupus erythematosus, rheumatoid arthritis. The present sequence represents a DNA encoding a PRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding PRO polypeptide, useful for diagnosing and treating an immune related disorder, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, thyroiditis, or diabetes mellitus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antirheumatic; Antiarthritic; Osteopathic; Hemostatic; Antianemic; Antithyroid; Antidiabetic; Nephrotropic; CNS-Gen.; Hepatotropic; Virucide; Gastrointestinal-Gen.; Antipsoriatic; Antiasthmatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADY14896 standard; protein; 174 AA
   satellite
                       Microsatellite instability colorectal tumor marker protein SEQ ID NO:53.
                                                  06-APR-2006
                                                                           AEF69875
                                                                                                 AEF69875 standard; protein;
                                                                                                                                                                                                                                                               Sequence 174
                                                                                                                                                                                                                                                                                     polypeptide
                                                                                                                                                                                                                                                                                                                                                                           Claim 8; SEQ ID NO 702; 158pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Abbas A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-AUG-2003; 2003US-0493546P
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                                                                                                                                                               FARYAR 74
                                                                                                                                                                                      FSRYAR 6
  DNA;
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                                                                                                                                                                                                               Conservative
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                                                  (first entry)
 tumor marker; therapeutic; diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ID NO 702.
                                                                                                                                                                                                                         90.3%;
                                                                                                                                                                                                                                                                                                   sequence represents a DNA encoding a PRO
                                                                                                                                                                                                             Score 28; DB Pred. No. 2.9e
1; Mismatches
                                                                                                   174 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Williams MP,
                                                                                                                                                                                                                         DB 9;
2.9e+02;
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prognosis,
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                                                                                                                                                                                                                                                                                                                                       antibody,
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RESULT 37
AEA17517
ID AEA17
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                          progression, or regression of cancer or a pre-malignant condition in a subject. for determining prognosis for cancer or a pre-malignant condition in a subject. The nucleotides and polypeptides are useful for determining the efficacy of a test compound or a therapy for inhibiting cancer in a subject and for selecting a composition for inhibiting cancer in a subject and for selecting a composition for inhibiting cancer in a subject. The compound, composition and method are useful for treating cancer. This sequence is a protein encoded by a marker DNA sequence specific for a high degree of microsatellite instability (MSI-H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting microsatellite instability, useful for monitoring the onset, progression, or regression of cancer, comprises measuring the expressi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 24; SEQ ID NO 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    colorectal tumor; cancer; neoplasm; cytostatic
                                             Thale cress caleosin-related family protein, SEQ ID NO:
                                                                               28-JUL-2005
                                                                                                                                                   AEA17517 standard; protein; 243
                                                                                                                                                                                                                                                                                                                                                                               Sequence 174 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                 marker) relative to colorectal tumors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUL-2004; 2004US-0591969P
13-APR-2005; 2005US-00105233
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MAYO FOUND MEDICAL EDUCATION & RES
                                                                                                                                                                                                                                                                                                         Similarity 5; Conserv
                                                                                                                                                                                                                                        FARYAR 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Catino TJ,
                                                                                                                                                                                                                                                                                                         Conservative
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                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                           90.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Burgart L,
                                                                                                                                                                                                                                                                                                         Score 28; DB 10; 1
Pred. No. 2.9e+02;
1; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                           Length 174;
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                                                                                                                                                                                                                                                                                                            Gaps
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Apolipoprotein; expression; transgenic plant; pharmaceutical; vascular disease; atherosclerosis; vasctropic; antiarterioscl

antiarteriosclerotic;

caleosin

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RESULT 38
ABR40670
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XXXXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                       CC pure oil bodies comprising an apolipoprotein obtained from plants. The curve oil bodies comprising an apolipoprotein obtained from plants. The curve of controlling expression is a seed-cc preferred or constitutive promoter such as a phaseolin promoter and a cc ubiquitin promoter. The chimeric nucleic acid construct additionally comprises a nucleic acid sequence encoding a stabilizing polypeptide cc linked in frame to the sequence encoding as stabilizing polypeptide cc specific oil body protein or a thioredoxin (SEQ ID NO: 56). The oil body protein is selected from oleosin, caleosin, and steroleosin. The non-cc plant specific stabilizing polypeptide is a green fluorescent protein cc (SEQ ID NO: 57) or a single chain antibody. The single-chain antibody is capable of facilitating purification of apolipoprotein expressed in seeds cor is capable of specifically associating with an oil body protein. Cc Alternatively, the single-chain antibody is a single-chain From A. Cc thaliana (D9scFv). The pharmaceutical apolipoprotein formulation may be considered to treat vascular diseases, such as atherosclerosis. The present
                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to methods for expressing apolipoprotein plants by introducing a chimeric nucleic acid construct into a plant cell. Specifically, this comprises providing a chimeric nucleic acid construct comprising a nucleic acid sequence capable of controlling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Expressing apolipoprotein in plants which may be used for treating vascular diseases, comprises introducing a chimeric nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-NOV-2003; 2003US-0519606P.
16-JUN-2004; 2004US-0579733P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell; and growing the plant cell into a mature plant where the plant expresses apolipoprotein. Also disclosed is a composition of substantially sure apolipoprotein obtained from a plant, or substantially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 122; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  construct into a plant cell.
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                 Zea mays oil trait related protein sequence SEQ ID NO:160
                                                      16-MAY-2003
                                                                                                                             ABR40670 standard; protein; 244
                                                                                                                                                                                                                                                                                                                                                                        Sequence 243
                                                                                                                                                                                                                                                                                                                                                                                                             sequence is a thale cress caleosin protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             expression in plant cells; and a sequence encoding an apolipoprotein polypeptide; introducing the chimeric nucleic acid construct into a p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Moloney MM, Reid A;
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                                                                                                                                                                                                                                                                                             Local Similarity res 5; Conserv
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                                                                                                                                                                                                                         FSKYAR 164
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                                                      (first entry)
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83.3%;
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Pred. No. 4.1e
1; Mismatches
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                                                                                                                                                                                                                                                                                          DB 9;
. 4.1e+02;
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                                                                                                                                                                                                                                                                                                                                     Length 243;
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RESULT 39
ADX91903
ID ADX91
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AC ADX91
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AC ADX91
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DT 21-AF
XX
XX
Plant
XX
XX
V
plant
XW
plant
XW
plant

ADX91903 standard; protein; 256

plant protectant; plant growth regulant; gene therapy; plant;
recombinant DNA construct; physical array; plant breeding marker;

Plant full length insert polypeptide seqid 54567.

(first entry

밁 Ś

Conservative

Indels

0

Gaps

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Query Match
Best Local Similarity
Thes 5; Conserv
                                                                                                                                     comprising a nucleic acid sequence (NS) chosen from a NS encoding a CC comprising a nucleic acid sequence (NS) chosen from a NS encoding a collapse potein (MAP)-kinase activity, LIPIS-like transcription factor activated protein (MAP)-kinase activity, LIPIS-like transcription factor activity, caleosin-like activity, ATP citrate lyase activity, SNF1-like activity, and CKC-like transcription factor activity. Also described: (1) complement (II) of (I); (2) a chimeric construct (III) comprising (I) or CC complement (II) in its genome; (4) sequence; (3) a plant (IV) and (5) collabsance from (V). (1) or its part can be used in antisense (2) inhibition or co-suppression in a transformed plant. (III) is useful for caltering the oil phenotype in a plant such as corn, soybean, wheat, rice, CC canola, Brassica, sorghum, sunflower or coconut. (III) is also useful for CC canola, Brassica, sorghum, sunflower or coconut. (III) is also be used as a hybridisation probe. ACC00626 to ACC00868 and ABR40591 to ABR40879 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Allen SM,
Jones TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plant; oil trait; oil phenotype; altered lipid profile; MAP kinase; receptor-like protein kinase; mitogen activated protein kinase; oil; LIP15-like transcription factor caleosin; ATP citrate lyase; SNF1; CKC-like transcription factor; antisense inhibition; co-suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel nucleotide fragment encoding polypeptides having receptor-like protein kinase activity, caleosin-like activity, useful for altering phenotypes in plants such as sunflower, coconut, soybean, wheat and r
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-JUN-2001; 2001US-0301913P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUN-2002; 2002WO-US020152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zea mays
                                                                                       Sequence 244 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 12; Page 202-203; 542pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB; ACC00705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-201509/19.
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                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes an isolated nucleotide fragment (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DUPO ) DU PONT DE NEMOURS & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Allen W
Kinney
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                                  90.3%;
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Klein TM,
                 Score 28; DB 6; Le
Pred. No. 4.2e+02;
1; Mismatches 0;
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Li C, Oliveira IC, Sakai
                                                     Length 244;
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ai H, Shen
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                                                                                         RESULT 40
                                                                                                                                                                                                                                                                                                  The invention describes a recombinant DNA construct comprising a construct comprising a consisting of a sequence encoding an amino acid sequence cavailable in electronic form from the US patent office at ftp.seqdata.uspto.gov/sequence.html?DocID:2004034888. The polynucleotide cof the invention are also useful in physical arrays of molecules and as comproving plant tolerance to cold, heat, drought, herbicides, extreme combination conditions, pathogens or pests, for manipulating growth rate in concreased resistance to plant disease, for manipulating growth rate in concreased resistance to plant disease, for producing galactonannan, compressed resistance to plant disease, for producing galactonannan, compressed resistance to plant disease, for increasing the rate of homologous recombination in plants, for improving yield by modification of photosynthesis or carbohydrate, nitrogen or phosphorus use and/or uptake corb by providing improved plant growth and development under at least one stress condition or for modifying seed oil or protein yield and/or content. This is the amino acid sequence of a plant full length insert of the invention.
                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cold tolerance; heat tolerance; drought tolerance; herbicide tolerance; extreme osmotic condition; pathogen tolerance; pest tolerance; growth rate; cell cycle pathway; disease resistance; galactomannan production; lignin production; plant growth regulator; yield; plant growth; plant development; seed oil; protein yield;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New recombinant DNA construct, useful for improving plant tolerance cold, heat, drought, herbicides, extreme osmotic conditions, pathoge pests, for conferring increased resistance to plant disease, or for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ZHOU/)
(KOVA/)
(SCRE/)
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 13-FEB-2002
                              ABG07163;
                                                       ABG07163 standard; protein; 301
                                                                                                                                                                                                                                                           Sequence 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 54567; 15pp; English
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05-NOV-2001;
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KOVALIC D K.
SCREEN S E.
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                                                                                                                                                                  FSRYAR 6
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(first
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entry)
                                                                                                                                                                                                             90.3%;
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Pred. No. 4.4e
1; Mismatches
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. 4.4e+02;
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or for
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RESULT 41
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and to produce other types of data and products dependent on DNA and amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated polynucleotide and encoded polypeptides, useful diagnostics, forensics, gene mapping, identification of mutat responsible for genetic disorders or other traits and to assemble to the contract of the contract o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymucleotides are also use in diagnostics as expressed sequence tags for identifying expressed in diagnostics as expressed sequence tags for identifying expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               biodiversity.
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23-AUG-2000; 2000US-00649167.
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                                                           28-JUL-2005
                                                                                                                                                                                   AEA17523 standard; protein; 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 20; SEQ ID NO 37522; 103pp;
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supplement; medical imaging; diagnostic; genetic disorder.
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                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 28; DB 4; Le
Pred. No. 5.2e+02;
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Thale cress

caleosin-1 protein,

SEQ ID

NO: 128

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RESULT 42
AEA17522
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                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Expressing apolipoprotein in plants which may be used for treating vascular diseases, comprises introducing a chimeric nucleic acid construct into a plant cell.
                                                                                                                                                                                                                                                                                                                                                               thaliana (D9scFv). The pharmaceutical apolipoprotein formulation may be used to treat vascular diseases, such as atherosclerosis. The present sequence is a thale cress caleosin protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 128; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2005-372355/38
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16-JUN-2004; 2004US-0579733P.
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                                                                                                                                                                                                                                                                                                                                  Sequence 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Moloney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-NOV-2004; 2004WO-CA001960
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28-JUL-2005
                                       AEA17522;
                                                                        AEA17522 standard; protein; 301
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                                                                                                                                                                                                                                                                                                                                  AA;
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(first entry)
                                                                                                                                                                                                                                                                        90.3%;
                                                                                                                                                                                                                                                   Score 28; DB 9;
Pred. No. 5.2e+02;
1; Mismatches 0
                                                                                                                                                                                                                                                                                          DB 9; Length 301;
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ABG27724
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AC ABG2
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                                                                            RESULT 43
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plants by introducing a chimeric nucleic acid construct into a plant cell. Specifically, this comprises providing a chimeric nucleic acid construct comprising a nucleic acid sequence capable of controlling expression in plant cells, and a sequence encoding an apolipoprotein polypeptide; introducing the chimeric nucleic acid construct into a plant cell; and growing the plant cell into a mature plant where the plant expresses apolipoprotein. Also disclosed is a composition of substantially pure apolipoprotein obtained from a plant, or substantially pure oil bodies comprising an apolipoprotein obtained from plants. The nucleic acid sequence capable of controlling expression is a seed-preferred or constitutive promoter such as a phaseolin promoter and a uniquitin promoter. The chimeric nucleic acid construct additionally constitutions.
                                                                                                                                                                                                                                                                                                                               comprises a nucleic acid sequence encoding a stabilizing polypeptide linked in frame to the sequence encoding apolipoprotein. It is a plant-specific oil body protein or a thioredoxin (SEQ ID NO: 56). The oil body protein is selected from oleosin, caleosin, and steroleosin. The non-plant specific stabilizing polypeptide is a green fluorescent protein (SEQ ID NO: 57) or a single chain antibody. The single-chain antibody is capable of facilitating purification of apolipoprotein expressed in seeds or is capable of specifically associating with an oil body protein. Alternatively, the single-chain antibody is a single-chain Fv antibody (SEQ ID NO: 240) capable of associating with the 18 kDa oleosin from A. thaliana (D9scFv). The pharmaceutical apolipoprotein formulation may be used to treat vascular diseases, such as atherosclerosis. The present sequence is a thale cress caleosin protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Expressing apolipoprotein in plants which may be used for treating vascular diseases, comprises introducing a chimeric nucleic acid construct into a plant cell.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to methods for expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 127; 530pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2005-372355/38
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16-JUN-2004; 2004US-0579733P.
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                                     ABG27724 standard; protein; 499
                                                                                                                                                                                                                                                                                              Sequence 301 AA;
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                                                                                                                                                                                                                 Conservative
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Pred. No. 5
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ABG27724;

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RESULT 44
ABG29681
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                                                                                                                                                                                                       Query Match
Best Local S
                                                                                                                                                                                              Matches
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23-AUG-2000; 2000US-00649167.
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   18-FEB-2002
                              ABG29681;
                                                        ABG29681 standard; protein; 520 AA.
                                                                                                                                                                                                                                                    Sequence 499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 20; SEQ ID NO 58083; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           biodiversity.
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                                                                                                                                    FARYAR 94
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                                                                                                                                                                                           Conservative
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 (first entry)
                                                                                                                                                                                                         90.3%;
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                                                                                                                                                                                         Score 28; DB 4; Le
Pred. No. 8.7e+02;
1; Mismatches 0;
                                                                                                                                                                                                                       Length 499;
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RESULT 45
ADN18185
ID ADN18
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AC ADN18
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DT 02-DE
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Bacterial polypeptide #838 02-DEC-2004 (first entry) ADN18185;

ADN18185 standard; protein; 681

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                                                                                                                                                                     CC genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is CC useful for generating antibodies against it, detecting or quantitating a CC uspplement. (II) and its binding partners are useful in medical imaging CC supplement. (II) and its binding partners are useful in medical imaging CC sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The CC diagnostics, forensics, gene mapping, identification of mutations or esponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and CC amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic control did not appear in the printed specification, but was obtained in CC electronic format directly from WIPO at CC fip.wipo.int/pub/published_pct_sequences
                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess biodiversity.
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23-AUG-2000; 2000US-00649167.
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                                                                                                                                              Sequence 520 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 20; SEQ ID NO 60040; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAS93868.
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supplement; medical imaging; diagnostic; genetic disorder.
46 FNRYAR 51
                                   1 FSRYAR 6
                                                                       5.
                                                                       Conservative
                                                                                          90.3%;
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                                                                                        Score 28; DB 4;
Pred. No. 9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New recombinant DNA construct comprising a promoter positioned to provide for expression of a polynucleotide encoding a polypeptide from a microbial source, useful for producing plants with improved properties.
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                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a recombinant DNA construct comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 838; 122pp; English.
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515
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) SLATER S C.
) CHEN X.
                                                                                                                                      from USPTO at segdata.uspto.gov/sequence.html.
                           1 FSRYAR 6
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                                                    Similarity 5; Conser
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FARYAR 520
                                                      Conservative
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                                                                   90.3%;
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Pred. No. 1
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                                                       Mismatches
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                                                                   DB 8;
1.2e+03;
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RESULT 46 ABG04036

ABP76771 ID ABP7 XX

ABP76771

standard; protein; 1206

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RESULT 47

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                                                                                                                                                                                        cc genes. (I) is useful in gene therapy techniques to restore normal cc activity of (II) or to treat disease states involving (II). (II) is consisted in tissue, as molecular weight markers and as a food cc polypeptide in tissue, as molecular weight markers and as a food cc supplement. (II) and its binding partners are useful in medical imaging consistes expressing (II). (I) and (II) are useful for treating disorders convolving abstrant protein expression or biological activity. The cc polypeptide and polynucleotide sequences have applications in convolving abstrant protein expression or biological activity. The cc polypeptide and polynucleotide sequences have applications in cd diagnostics, forensics, gene mapping, identification of mutations cc and no produce other types of data and products dependent on DNA and canno acid sequences. ABG00010-ABG30377 represent novel human diagnostic camino acid sequences of the invention. Note: The sequence data for this captain in the printed specification, but was obtained in celectronic format directly from WIPO at
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Matches
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23-AUG-2000; 2000US-00649167.
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                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 20; SEQ ID NO 34395; 103pp; English.
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                                                                                  Local
                                                                   Similarity 5; Conser
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                                FSRYAR 6
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                                                                   Conservative
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                                                                 Score 28; DB 4;
pred. No. 1.5e+03;
1; Mismatches 0
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                                                                                                       Length 847;
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RESULT 48
ABP78681
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                                                                                                                                                                                                                                                                          Matches
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Best Local
                                                                                                                                                                                                                                                                                                                       antibodies that specifically bind to the proteins. The composition comprising the protein, nucleic acid molecules encoding the proteins and comprising the protein, nucleic acid or antibody is useful for the manufacture of a medicament for treating or preventing N. gonorrhoeae infection, this may be in the form of a vaccine or gene therapy. Sequences given in records ABP76736-ABP81046 represent nucleic acid molecules of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-MAR-2003
                 12-FEB-2001; 2001GB-00003424.
                                    12-FEB-2002;
                                                       10-OCT-2002
                                                                         WO200279243-A2
                                                                                           Neisseria gonorrhoeae
                                                                                                              Antibacterial;
                                                                                                                                                   07-MAR-2003
                                                                                                                                                                      ABP78681
                                                                                                                                                                                     ABP78681 standard; protein; 1206 AA
                                                                                                                                                                                                                                                                                                              Sequence 1206 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 182; 815pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    New protein from Neisseria medicament for treating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fontana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-FEB-2002; 2002WO-IB002069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neisseria gonorrhoeae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antibacterial; infection; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABP76771;
(CHIR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHIR-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gonorrhoeae amino acid sequence
                                                                                                                               gonorrhoeae
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DB; ABZ37741.
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CHIRON
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                                    2002WO-IB002069
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                                                                                                                                                  (first entry)
SPA
                                                                                                              infection; vaccine;
                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                  90.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Masignani
                                                                                                                              sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                   gonorrheae,
preventing |
                                                                                                                                                                                                                                                                                  Score 28;
Pred. No.
                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                               SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                    N. gonorrheae infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                              useful for the manufacture of
                                                                                                                                                                                                                                                                                   2.1e+03;
                                                                                                                                ID 3892.
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RESULT 49
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to proteins from Neisseria gonorrhoeae. Also disclosed are the nucleic acid molecules encoding the proteins ar antibodies that specifically bind to the proteins. The composition comprising the protein, nucleic acid or antibody is useful for the manufacture of a medicament for treating or preventing N. gonorrhoeae infection, this may be in the form of a vaccine or gene therapy. Sequences given in records ABP76736-ABP81046 represent nucleic acid molecules of the invention
The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of
                                                                                                             Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from bacterial infection.
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                                                                                                                                                                                                     Rubenfield MJ,
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27-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                  Pseudomonas aeruginosa
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                                                                                                                                                                                                                                                                                            18-FEB-1999;
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                                                                                                                                                                                                                              (GENO-) GENOME
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5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; protein; 1778 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                       SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                              THERAPEUTICS CORP.
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98US-0094190P.
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                                                                                       18159;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gonorrheae, useful preventing N. gonor
                                                                                                                                                                                                     Deloughery
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 28;
Pred. No.
                                                                                     455pp;
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                                                                                      English.
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2.1e+03;
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                                                                                                                                                                                                      Bush
                                                                                                                                                                                                                                                                                                                                                                                              infection; antibacterial.
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                                                                                                  The present sequence represents a zebrafish alpha1C subunit of a voltage-dependent I-type calcium channel. The specification describes a method for determining whether a test subject has or is at risk of developing a disease or condition related to the alpha1C subunit. The method involves analysing a nucleic acid molecule sample from the test subject to determine whether the test subject has a mutation in a gene encoding the alpha1C subunit, where the presence of a mutation indicates that subject has or is at risk of developing the disease. The method is useful for determining whether a test subject (preferably mammal e.g. human) has, or is at risk of developing, a disease or condition related to the alpha1C, is at risk of developing, a disease or condition related to the alpha1C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Determining whether a subject has or is at risk of developing disease related to alphalC subunit of voltage-dependent L-type calcium channel, by detecting mutations in genes encoding the subunit in sample of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-FEB-2002; 2002WO-US003441.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 57-61; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AlphalC subunit of a voltage-dependent L-type calcium channel.
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   Sequence 2151 AA;
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                                                                            t risk of developing, a deart disease including
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pred. No. 3.2e+03;
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                                                                            the alphalC, fibrillation
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Pred. No. 3.9e+03;
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1 31 3 31 3 31 4 3 31 6 3 31 7 3 31 7 3 31 10 28 11 2 28 11 2 28 11 2 28 12 28 14 28 12 28 14 28 15 28 16 28 17 28 18 28 19 28 27 27 28 27 29 27 29 27 29 27 29 27	Result No. Score		Pred. No. score gree and is de	Database :	Post-processing:	Minimum DB seq Maximum DB seq	Total number of	Searched:	Scoring table:	Title: Perfect score: Sequence:	Run on:	OM protein - pr	
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C64890 C72022 F8603 C69997 F86051 D91075 C85920 C85920 T16828 T16828 T16828 T16828 C909202 E64908 C90897 D85720 D8		SUMMARIES	Pred. No. is the number of results predicted by chance to h score greater than or equal to the score of the result bein and is derived by analysis of the total score distribution.		0% 100% 500 summaries		parameters:	63 residues	0.5		06:00:58 ; Search time 6 Seconds (without alignments) 96.217 Million cell updates/sec	sw model	e version 5.1.9 3 - 2006 Biocceleration
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RESULT 2 C72022 UDP-N-acetylglucosamine-N-acetylmuramyl- (pentapeptide)	Qy 1 FSRYAR 6 Db 80 FSRYAR 85	Query Match 100.0%; Score 31; DB 2; I Best Local Similarity 100.0%; Pred. No. 14; Matches 6; Conservative 0; Mismatches 0;	ron transfer; iron-sulfur pr mbrane #status predicted <tm xxin [2Fe-2S] homology <fer> y site: 2Fe-2S cluster (Cys)</fer></tm 	;Status: nuclear acid sequence not snown; trann ;Molecule type: DNA ;Residues: 1-356 <blat> ;Cross-references: UNIPROT:P76081; UNIPARC:UPII; ;Experimental source: strain K-12, substrain M(;Experimental source: strain K-12)</blat>	J.; Mau, B.; Shao, Y. 1453-1462, 1997 complete genome sequence of Es umber: A64720; MUID:97426617; C64890	nsfer prote .a coli !sequence_re	ALIGNMENTS	97 23 74.2 363 2 98 23 74.2 366 2 99 23 74.2 366 2 99 23 74.2 366 2 00 23 74.2 367 2	23 74.2 360 2 23 74.2 360 2 23 74.2 360 2 23 74.2 361 1 23 74.2 362 2	86 23 74.2 360 1 87 23 74.2 360 2 88 23 74.2 360 2 89 23 74.2 360 2 90 23 74.2 360 2 91 23 74.2 360 2	81 23 74.2 358 2 82 23 74.2 358 2 83 23 74.2 358 2 84 23 74.2 358 2 85 23 74.2 359 2 85 23 74.2 359 2	470 23 74.2 347 2 875)12 471 23 74.2 348 2 B70109 472 23 74.2 348 2 B83869 473 23 74.2 348 2 B96820 474 23 74.2 348 2 I51677 475 23 74.2 348 2 I51677 476 23 74.2 350 2 AH3020 477 23 74.2 350 2 AP302116 478 23 74.2 350 2 JC2116 479 23 74.2 351 2 JC2116 479 23 74.2 352 2 JC2116 479 23 74.2 352 2 JC216	68 23 74.2 346 2 69 23 74.2 346 2
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R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzber Nucleic Acids Res. 28, 1397-1406, 2000

A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.

A;Reference number: A81500; MUID:20150255; PMID:10684935
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A;Cross-references: UNIPROT:Q9Z702; UNIPARC:UPI00000470A7; GB:AE001670; GB:AE001363; NII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptidoglycan transferase [imported] - Chlamydophila pneumoniae (strain J138)
C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae
C;Date: 02-Mar_2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
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C;Superfamily: murG
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                       probable proline transport protein - Bacillus subtilis
N;Alternate names: probable proline-specific permease
C;Species: Bacillus subtilis
                                                                                                           C69997
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A;Title: Comparison of whole genome sequences of chlamydia pneumoniae J138.
A;Reference number: A86491; MUID:20330349; PMID:10871362
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A;Experimental source: strain AR39, HL cells
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C; Species: Bacillus C; Date: 05-Dec-1997
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#sequence_revision 05-Dec-1997 #text_change 05-Oct-2004
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                                                                                                                                                                                                                                                                                                                                                              Science 277, 1453-1462, 1997
A,Title: The complete genome sequence of Escherichia coli K-12
A,Reference number: A64720; MUID:97426617; PMID:9278503
                                                                              Mol. Biol. Evol. 9, 688-706, 1992
A; Title: Nucleotide sequence, function, activation, and number: A44070; MUID:92334140; PMID:1630307
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                                                                                                                                                                                                      A;Cross-references: UNIPROT:P24240;
A;Experimental source: strain K-12;
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A; Residues: 1-474 <BLAT>
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source: strain 9, 688-706, 1992

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GB:AE000355;

GB:U00096; NID:92

evolution

of the cryptic asc

ope:

acid sequence not

shown; translation

not

Burland, V.; Riley,

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R;Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Bertert C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Choi A.; Ehrlich, S.D.; Emmerson, P.T.; Ehrlian, K.D.; Errington, J.; Fabret, C.; Ferrari, E. Nature 390, 249-256, 1997

A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallert iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F.; Koetter, P.; Koningstein, G.; Krogth, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, Koetter, P.; Koningstein, G.; Krogth, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, M.; Jogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Sato, T.; Scanlon, R; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sadaie, Y.; Sato, T.; Scanlon, A; Mitters, P.; Wijat, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama, T.; Winters, P.; Wijat, A.; Yamamoto, H.; Vamanoto, K.; Yata, K.; Yoshida, K., A; Hiters, P.; Wijat, A.; Yamanoto, H.; Vamanoto, K.; Yata, K.; Yoshida, K., A; Ference number: A69580; MUID:98044033; PMID:9384377
A; Accession: C6997
A, McCession: C6997
A, McCession: Sreliminary; nucleic acid sequence not shown; translation not shown
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A;Residues: 1-463 <KUN>
A;Cross-references: UNIPROT:034618; UNIPARC:UPI0000060942; GB:Z99119; GB:AL009126; NID:g2
A;Experimental source: strain 168
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Best Local
   76
                                                              1 FSRYAR
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   FSRYAR
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   81
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Pred. No.
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6-phospho-beta-glucosidase [imported] - Escherichia coli (strain 0157:H7, substrain C;Species: Escherichia coli (c;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004 C;Accession: D91075 R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H. DNA Res. 8, 11-22, 2001 A;Ritle: Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and A;Reference number: A99629; MUID:21156231; PMID:11258796
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A;Cross-references: UNIPARC:UPI000016EE0A; GB:M73326; NID:g145385; PIDN:AAA16430.1;
A;Experimental source: strain LP103
A;Experimental source extracted from NCBI backbone (NCBIN:109109, NCBIP:109114)
C;Genetics:
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C;Superfamily: Agrobacterium beta-glucosidase
C;Keywords: glycosidase; hydrolase
                                                                                                                                                          A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-474 <STO>
                                                                                                                                                                                                                                                                   R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Gene: ECs3572
C;Superfamily: Agrobacterium beta-glucosidase
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                                                                                                                                                                                                                                                                                                                                                          6-phospho-beta-glucosidase, cryptic [imported] - Escherichia coli (strain O157:H7, C;Species: Escherichia coli
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A;Experimental source: strain O157:H7, substrain RIMD 0509952
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A; Residues: 1-474 <HAY>
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hypothetical protein T07F12.1 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #te: C;Accession: T16828 R;Chissoe, S.
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C;Superfamily:
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A; Residues: 1-496 <STO>
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A;Molecule type: mRNA
A;Residues: 1-239 <HSU>
A;Cross-references: UNIPROT:023959; UNIPARC:UPI00000A8419; EMBL:AF004809; NID:g2270993; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ca+2-binding EF hand protein homolog PM13 - soybean C;Species: Glycine max (soybean) C;Species: Glycine max (soybean) C;Date: 30-Apr.1999 #sequence_revision 30-Apr.1999 #text_change 09-Jul-2004 C;Accession: T07092 .....
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R;Stover, C:K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, ar
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C;Superfamily: rice abscisic acid-induced protein
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A;Reference number: A82950; MUID:20437337; PMID:10984043
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C;Species: Pseudomonas aeruginosa
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Pred. No. 19;
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    Pseudomonas aeruginosa

20-Sep-1999 #text_change 09-Jul-2004
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K.; Lim,
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C;Accession: AF3271
R;DelVecchio, V.G., Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Ivanova, R;DelVecchio, V.G., Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.; Ivanova, P.; Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan, D.; Letess Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
A;Accession: AF3271
                                  A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-437 < KUR>
A;Cross-references: UNIPROT:Q92X34; UNIPARC:UPI00000CB40E; GB:AL591985; PIDN:CAC48528.1;
A;Cross-references: UNIPROT:Q92X34; UNIPARC:UPI00000CB40E; GB:AL591985; PIDN:CAC48528.1;
A;Experimental source: strain 1021, megaplasmid pSymB
A;Experimental source: strain 
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A;Molecule type: DNA
A;Residues: 1-409 <KUR>
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A;Molecule type: DNA
A;Residues: 1-304 <CHI>A;Cross-references: UNIPROT:Q22323; UNIPARC:UPI000007A68E; EMBL:U29154; NID:g861403;
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A;Title: The complete sequence of the 1,683-kb pSymB megaplasmid from the N2-fixing enda A;Reference number: A95842; MUID:21396508; PMID:11481431
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C;Species: Sinorhizobium meliloti
C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 31-Dec-2004
C;Accession: H95857
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A;Experimental source: strain 16M
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A;Authors: Kahn, D.; Kahn, M.L.; Kalman,
                                                                                                                                                                                                                                                                                                                                                                                                       A; Accession: H95857
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;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 09-Jul-2004
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        A;Status: preliminary A;Molecule type: DNA
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methionyl-tRNA synthetase (metS) [imported] - Sulfolobus solfataricus C;Species: Sulfolobus solfataricus C;Species: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004 C;Accession: D99202 C;Riches, C.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.; Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001 A;Description: Sulfolobus solfataricus complete genome. A;Accession: D90202 A;Accession: D90202
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A;Residues: 1-475 <KUR>
A;Cross·references: UNIFROT:Q92YG7; UNIPARC:UPI00000CB248; GB:AE006469; PIDN:AAK65571.1;
A;Cross·references: UNIFROT:Q92YG7; UNIPARC:UPI00000CB248; GB:AE006469; PIDN:AAK65571.1;
A;Cross·references: UNIFROT:Q92YG7; UNIPARC:UPI00000CB248; GB:AE006469; PIDN:AAK65571.1;
A;Cross·references: UNIFROT:Q92YG7; UNIPARC:UPI0000CB248; GB:AE006469; PIDN:AAK65571.1;
A;Cross·references: UNIFROT:Q92YG7; UNIPARC:UPI0000CB248; GB:AE006469; PIDN:AAK65571.1;
A;Cross·references: LAFOUNCES: Replay CB:AE006469; PIDN:AAK65571.1;
A;Cross·references: LAFOUNCES: Left Composite Genome of the Legume Symbiont Sinorhizobium meliloti.
A;Reference number: A96039; MUID:21368234; PMID:111474104
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A;Title: Nucleotide sequence and predicted functions of the entire Sinorhizobium meliloti A;Reference number: A95262; MUID:21396509; PMID:11481432
A;Accession: A95376
A;Status: preliminary
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C;Superfamily: ecotropic
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A;Residues: 1-573 <1
A;Cross-references:
C;Genetics:
dipeptidyl carboxypeptidase II [imported] - Escheri
C;Species: Escherichia coli
C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001
C;Accession: C90897
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A; Residues: 1-138'LL',141-681 <HEN>
A; Cross-references: UNIPARC: UPI000016F0BD;
A; Becker, S.; Plapp, R.
submitted to the EMBL Data Library, Februar
A; Reference number: S14870
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                                                                                                                                              RESULT
C90897
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C;Superfamily: methionyl-tRNA synthetase
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A;Cross-references: UNIPARC:UPI000016F0BD;
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A; Residues: 1-681 <BLAT>
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Pred. No. 1.3e+02;
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                             A; MOLTECUARY (1877) - KURY
A; Cross-references: UNIPROT: 08YS49; UNIPROT: 08YS49; UNIPROT: 7120
                                                                                                                                                                                                                                                                                                              C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change C;Accession: AC2211
C;Accession: Accession: A
   A; Experimental source: C; Genetics:
                                                                                                                                                                                                            DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nit A;Reference number: AB1807; MUID:21595285; PMID:11759840 A;Accession: AC2211
                                                                                                                                       A; Molecule type:
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Nature 409, 529-533, 2001
A;Title: Genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A85480; MUID:21074935; PMID:11206551
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C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 09-Jul-2004
C;Accession: D85720
C;Accession: D85720
C;Accession: D15720
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A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Accession: C90897
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C;Superfamily:
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C;Species:
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A; Residues: 1-681 <H
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UNIPARC: UPI00000CE703;

GB:BA000019; PIDN:BAB74941.1;

Filamentous Nitrogen-fixing

Cyanobacterium

Anat

S.; Watanabe, , M.; Yasuda, I

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Tabata, Iriguchi, Tabata, S

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strain PCC 09-Jul-2004

(strain PCC

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A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana. A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: F84791
                                                                                                                                                                                                                                                                                                                                                                                                                  R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Fujii, C.Y.; M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; Vanaken, S.E.; Umayam, L.; Tallon, L. euss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, C. Nature 402, 761-768, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, A;Reference number: A82950; MUID:20437337; PMID:10984043 A;Accession: D83181
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Nature 406, 959-964, 2000
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A; Residues: 1-36 <STO>
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A;Residues: 1-1746 <STO>
A;Cross-references: UNIPROT:Q9HXR4; UNIPARC:UPI00000C5AE2;
A;Experimental source: strain PAO1
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AI0982
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A.; Larbig,
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K.; Lim,
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Nature 413, 848-852, 2001

A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; A;Title: Complete genome sequence of a multiple drug resistant A;Reference number: AB0502; MUID:21534947; PMID:11677608

A;Accession: AI0982
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VCO33 protein homolog (AF179595) [imported] - Agrobacterium tumefaciens (strain C58, C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004 C;Accession: F97521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Parkhill, J.; Dougan, G.; James, K.D.; Thomson, N.R.; Pickard, D.; Wain, J.; Cth, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Dowd, L.; White, N.;
                                                         Karp, P.; Romero, P.; Zhang,
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Bi
ster, E.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Quro. A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, Science 294, 2323-2328, 2001
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C;Genetics:
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A; Residues: 1-43 < PAR>
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A; Title: The Genome A; Reference number: A; Accession: AH2740
                                                                                                                              R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, erage, G.; Gillet, W.; Grant, C.; Guenthner, ; Karp, P.; Romero, P.; Zhang, S.
                                                                                                                                                                                 conserved hypothetical protein Atul336 [imported] - Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens C;Decies: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004 C;Accession: AH2740
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A;Reference number: A97359; MUID:21608551; PMID:11743194
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                      of the Natural Genetic Engineer Agrobacterium tumefaciens C58 AB2577; MUID:21608550; PMID:11743193
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83.3%;
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                                                                                    W.; Perry, M.; Gordon-Kamm,
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kelz, B.;
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A;Gene: guk
A;Introns: 25/1; 80/3; 83/1; 173/2
A;Introns: 25/1; 80/3; 83/1; 173/2
C;Superfamily: mucleoside diphosphate kinase
C;Keywords: ATP binding; hexamer; mitochondrion; phosphohistidine; phosphoprotein;
C;Keywords: ATP binding #status predicted
F;83-87/Region: ATP binding #status predicted
F;186/Active site: His (phosphohistidine intermediate) #status predicted
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A;Molecule type: DNA
A;Residues: 1-108 <KUR>
A;Cross-references: UNIPROT:QBUFR1; UNIPARC:UPI00000D1B08; GB:AE008688; PIDN:AAL42342.1.
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A; Residues: 1-220 < TRO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Status: preliminary
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A; Residues: 1-208 <STO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
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Matches 5
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                                 Query Match
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   Similarity
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Pred. No.
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Score 27;
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                                                                                                                                                                                                                                                                                                                                                                  GB:L23068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PMID:8244981
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A; Decirce type: DNA
A;Molecule type: DNA
A;Residues: 1-69,'I',71-220,'A' <WOO>
A;Cross-references: UNIPARC:UPI0000133CD5; EMBL:AL035077; PIDN:CAA22664.1; GSPDB:GN00066;
                                                                                                                                                                                                                                                                                                        R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, N.; Karp, P.; Romero, P.; Zhang, S.; Science 294, 2317-2323, 2001
                                                                                                                                                                                                                                                                                                                                                                                                 C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004 C;Accession: AE3245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Map position:
C; Superfamily: 1
C; Keywords: cyto
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A;Reference number: Z21878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Masson, J.Y.; Vadnais, J.; Ramotar, D. Gene 170, 153-154, 1996
A;Title: The Schizosaccharomyces pombe spqM gene is a new A;Reference number: JC4755; MUID:96200877; PMID:8621081
A;Accession: JC4755
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C;Date: 10-May-1996 #sequence_revision 16-Aug-1996
C;Accession: JC4755; T39755
A; Genome: plasmid
C; Superfamily: ol:
                                                                                    A;Cross-references: UNIPROT:Q8U604;
A;Experimental source: strain C58 ()
                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-286 < KUR>
                                                                                                                                                                                                  A;Reference number:
A;Accession: AE3245
                                                                                                                                                                                                                      A; Title: The Genome of the Natural Genetic Engineer Agrobacterium A; Reference number: AB2577; MUID:21608550; PMID:11743193
                                                                                                                                                                                                                                                               A; Authors: Yoo, H.; Tao, ster, E.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein accE [imported] - C;Species: Agrobacterium tumefaciens
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A; Residues: 1-232 < MAS>
                                           A;Gene: accE
                                                                  C; Genetics:
                                                                                                                                                                           A; Status: preliminary
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83.3%;
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Pred. No.
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                                                                                      ; UNIPARC:UPI00000D27AF; (Dupont)
                                                                                                                                                                                                                                                                                      P.; Jung, M.; Krespan,
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Local Similarity

87.1%; 83.3%;

Score Pred.

27; No.

DB 2;

Length

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C;ACCEBBLUM.
R;Liao, M.K.; Maloy, S.
submitted to the EMBL Data Library,
submitted To The EMBL Data Library,
Tdentification of a
          A;Description: Identification
A;Reference number: S35983
A;Accession: S35983
                                                                                                                                                                                                                                                                                                                                                                                                                         proline transport protein - Salmonella typhimurium C;Species; Salmonella typhimurium C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995
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A; Residues: 1-286 < P.
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A; Accession: T03434
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C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999
C;Accession: T03434
R;Piper, K.R.; Beck von Bodman, S.; Cook, D.M.; Hwa
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C;Superfamily: oligopeptide permease protein
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A; Reference number: A87249; MUID:21173698; A; Accession: E87697
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A;Residues: 1-292 <LIA>
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83.3%;
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A;Reference number: Z23016
A;Reference number: Z23016
A;Accession: T45890
A;Status: preliminary
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-302 <BLO>
A;Cross-references: UNIPROT:Q9FYB7; UNIPARC:UPI000017A395;
A;Cross-references: cultivar Columbia; BAC clone F4P12
                                                                                                                      R.Dairi. T., Hamano, Y.; Igarashi, Y.; Furumai, T.; Oki, T.

Biosci. Biotechnol. Biochem. 61, 1445-1453, 1997
A;Title: Cloning and nucleotide sequence of the putative polyketide synthase genes for A;Reference number: JC5850; MUID:97480928; PMID:9339544
A;Accession: JC5855
A;Molecule type: DNA
A;Residues: 1-341 <DAI>
A;Cross-references: UNIPROT:032456; UNIPARC:UPI000002F07D; DDBJ:D87924; NID:g2580441; C;Comment: This enzyme catalyzes repeated condensation cycles of acyl-CoA, affecting the condensation.
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A;Gene: CC3615
C;Superfamily: dTDP-dihydrostreptose synthase
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N;Alternate names: protein F4P12.200
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change
C;Accession: T45890
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A; Residues: 1-293 <STO>
                                                                                                                                                                                                                                                                                                             polyketide synthase (EC 2.-.-.) chain 6 - Actinomadura hibisca C;Species: Actinomadura hibisca C;Date: 05-Mar-1998 #sequence_revision 13-Mar-1998 #text_change C;Accession: JC5855
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A; Introns: 36/2; 41/3; 67/3; 99/3; 164/2; 198/2
A; Note: F4P12.200
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Matches 5
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M.; Ohfu
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A; Title:
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C;Species: Pyrococcus abyssi
C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999
C;Accession: E75080
R;anonymous, Genoscope
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                                                               probable phosphoglycerate kinase - Pyrococcus horikoshii
C;Species: Pyrococcus horikoshii
C;Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text_change 16-Aug-2004
C;Accession: D71065
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              R;Kawarabayasi, Y.; Sawada, M.; Ohfuku, Y.; Funahashi, DNA Res. 5, 55-76, 1998
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C; Superfamily:
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A;Accession: E75080
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C;Superfamily:
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A; Residues: 1-354 <STO>
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                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-410 < KAW>
                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
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C;Species: Pseudomonas aeruginosa
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Experimental source: strain PAO1
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T.; Tanaka, T.; Kudoh,
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J.; Kushida,
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A.; Larbig,
hyper-thermophilic
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K.; Lim,
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                                                Sekin
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                                                A;Gene:
C;Superf
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D87672
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                                                                                             A; Cross-references:
                                                                                                               A;Residues:
                                                                                                                            A; Molecule type: DNA
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Best Local
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Superfamily:
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 Similarity
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A,Status: preliminary; nucleic acid sequence not shown; tra
A,Molecule type: DNA
A;Residues: 1-410 <KAW>
A,Cross-references: UNIPROT:058965; UNIPARC:UPI000013180B;
A,Experimental source: strain OT3
A,Note: this accession replaces an interim accession for a
C;Genetics:
A,Gene: PH1218
C;Superfamily: phosphoglycerate kinase
                                                                                                                                                                                                                                                               R,Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, J.I. B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolong, J.; Ermolaeva, Mitte, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Experimental source: strain C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            submitted to the EMBL Data Library, July 1995 A; Description: sequence of C. elegans cosmid A; Reference number: S59417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein C30B5.3 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #t0;Accession: T15705
                                                                                                                                                                                                                        A;Reference number: A87249; MUID:21173698; PMID:11259647 A;Accession: D87672
                                                                                                                                                                                                                                                                                                                                                                               C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change 05-Oct-2004
C;Accession: D87672
                                                                                                                                                                                                                                                                                                                                                                                                                           glutamate-cysteine ligase [imported] -
C;Species: Caulobacter crescentus
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A;Introns: 12/2; 91/1; 349/3
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A; Residues: 1-411 < DUZ >
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                                                                                                                                                                                                   A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPROT:Q18317; UNIPARC:UPI0000075091; EMBL:U23450; NID:g733552; PID:
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                                                                                                                                  UNIPROT:Q9A2Z2; UNIPARC:UPI00000C7A81; GB:AE005673; NID:g13425126; PJ
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C;Species: Escherichia coli
C;Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_cl
C;Accession: B64769
R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.'
A.; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli
A;Reference number: A64720; MUID:97426617; PMID:9278503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:P77327; UNIPARC:UPI000013232F; A;Experimental source: strain K-12, substrain MG1655 C;Superfamily: ecotropic retrovirus receptor protein C;Keywords: amino acid transport; transmembrane protein F;43-59/Domain: transmembrane #status predicted <TM1> F;96-112/Domain: transmembrane #status predicted <TM2>
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Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant
A;Reference number: AB0502; MUID:21534947; PMID:11677608
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C; Superfamily:
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A; Residues: 1-456 < PAR>
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Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 05-Oct-2004;
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              Similarity 5; Conserv
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Cronin, A.;
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              probable proline-specific permease C;Species: Yersinia pestis C;Date: 02-Nov-2001 #sequence_revis C;Accession: A10388 C;Parkhill, J.; Wren, B.W.; Thomson
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proline permease transport protein ECs0452 [imported] - Escherichia coli (strain O157:H7, C;Species: Escherichia coli (C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 05-Oct-2004 C;Accession: D90685 C;Accession: D90685 R;Hayysshi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; Han, C.G.;
                                                                                                                                                                                                                                                                                                                                                A;Molecule type: DNA A;Residues: 1-457 <STO> A;Cross-references: UNIPROT:P77327; UNIPARC:UPI000013232F; A;Experimental source: strain O157:H7, substrain EDL933 C;Genetics:
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C; Superfamily:
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Nature 409, 529-533, 2001
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C;Superfamily:
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A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A:Reference number: A99629; MUID:21156231; PMID:11258796
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DNA Res. 8, 11-22, 2001
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A;Accession: D90685
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Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; S
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#sequence_revision 02-Nov-2001 #text_change

[imported] - Yersinia

pestis (strain CO92)

05-Oct-2004

B.W.; Thomson,

N.R.; Titball,

R.W.; Holden, M.T.G.; Prentice,

м. В.;

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phospho-beta-glucosidase ArbB - Erwinia chrysanthemi
C;Species: Erwinia chrysanthemi
C;Date: 04-Sep-1998 #sequence_revision 04-Sep-1998 #text_change 09-Jul-2004
C;Accession: C42603
R;el Hassouni, M.; Henriasat D. C.'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;el Hassouni, M.; Henrissat, B.; Chippaux, M.; Barras, F.
J. Bacteriol. 174, 765-777, 1992
A;Title: Nucleotide sequences of the arb genes, which control beta-glucoside utilization clase family including enzymes from eubacteria, archeabacteria, and humans.
A;Reference number: A42603; MUID:92121114; PMID:1732212
A;Accession: C42603
                                                                                                                                                                                                                         R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, J. B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolon n, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
                                                                                                                                                                                                                                                                                                                                                    C;Date: 20-Ap
C;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPROT:P26206; UNIPARC:UPI0000175B1C A;Note: sequence inconsistent with the nucleotide translation A;Note: sequence extracted from NCBI backbone (NCBIN:77535, NCC;Superfamily: Agrobacterium beta-glucosidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-465 <EL1>
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A; Gene: CC0822
C; Superfamily: NAD-dependent aldehyde dehydrogenase; aldehyde dehydrogenase homology
                                                                                                                                                                                                                                                                                                                                                                                                     aldehyde dehydrogenase [imported] - Caulobacter crescentus
C;Species: Caulobacter crescentus
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C;Superfamily:
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A; Residues: 1-463 < KUR>
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                                                                            A;Cross-references: UNIPROT:Q9A9Y9; UNIPARC:UPI00000C71A2; GB:AE005673; NID:g13422073;
                                                                                                         A; Residues: 1-478 <STO>
                                                                                                                          A; Molecule type: DNA
                                                                                                                                                       A;Status: preliminary
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                                                                                                                                                                                                       Reference number: A87249;
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5; Conserv
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83.3%;
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Pred. No. 1.5e+02;
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A;Cross-references: UNIPROT:Q9Z7V7; UNIPARC:UPI0000043CB3; GB:AE001644
A;Experimental source: strain CWLO29
R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; W
C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McCla
Nucleic Acids Res. 28, 1397-1406, 2000
A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia
A;Reference number: A81500; MUID:20150255; PMID:10684935
                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Kalman, S.; Mitchell, W.; Marathe, R.; Lammel, C.; Fan, Canture Genet. 21, 385-389, 1999
A;Title: Comparative genomes of Clamydia pneumoniae and C. A;Reference number: A72000; MUID:99206606; PMID:10192388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide ABC transporter, permease protein, probable CP0151 [imported] - Chlamydophila C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004 C;Accession: C72059; B81608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Shirai, M.; Hirakawa, H.; Kimoto, M.; Tabuchi, M.; Kishi, Nucleic Acids Res. 28, 2311-2314, 2000
A;Title: Comparison of whole genome sequences of chlamydia A;Reference number: A86491; MUID:20330349; PMID:10871362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligopeptide permease [imported] - Chlamydophila pneumoniae (strain J138) C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae (C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004 C;Accession: B86565
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                                                                                                                                   A; Experimental source: strain C; Genetics:
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A; Residues: 1-579 < REA>
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A; Residues: 1-579 < ARN>
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                                   Similarity 5; Conserv
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Pred. No. 1
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Pred.
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1.9e+02;
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FSRYVR

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A;Gene: nodM
A;Genome: plasmid pRLIJI
C;Superfamily: glutamine-f
C;Keywords: aminotransfera
F;2-608/Product: glutamine
F;2/Active site: Cys #stat
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C; Superf
C; Keywor
glutamine-fructose-6-phosphate transaminase N;Alternate names: glucosamine-fructose-6-p C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision 30-Se C;Accession: C97575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Accession: $01040
R;Surin, B.P.; Downle, J.A.
Mol. Microbiol. 2, 173-183, 1988
A;Title: Characterization of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           glutamine-fructose-6-phosphate transaminase (isomerizing)
N;Alternate names: nodulation protein nodW
C;Species: Rhizobium leguminosarum bv. viciae
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_c
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C;Accession: S16561
C;Accession: S16561
C;Accession: S16561
Petrovics, G.; Banfalvi, Z.; Kondorosi, A.
Mol. Gen. Genet. 228, 113-124, 1991
Mol. Gen. Genet. 228, 113-124, 1991
A;Title: Six nodulation genes of nod box locus 4 in Rhizobium meliloti
A;Reference number: S16561; MUID:91360053; PMID:1909418
A;Accession: S16561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Rhizobium meliloti
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995
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A; Molecule type: DNA
A; Residues: 1-608 <SUR>
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A; Residues: 1-605 <BAE>
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                                                                                                                                                                                                                                                                                                                       Superfamily: glutamine-fructose-6-phosphate aminotransferase (isomerizing); Reywords: aminotransferase; isomerase; nodulation; 2-608/Product: glutamine-fructose-6-phosphate transaminase (isomerizing) #status; 2/Active site: Cys #status predicted
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2/Active site: Cys #status predicted
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Pred. No. 2e+0
0; Mismatches
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Pred. No. 2e+0
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PMID:3132583
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-608 «KUR»
A;Cross-references: UNIPROT:Q8UEH1; UN
A;Cross-references: Strain C58 (Dug
C;Genetics:
A;Gene: glmS
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A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrob A;Reference number: A97359; MUID:21608551; PMID:11743194
A;Status: preliminary
                                                                                                                                                                                                                                                                                                     ster, E.W.
A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58
A;Reference number: AB2577; MUID:21608550; PMID:11743193
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A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein glms [imported] - C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision C;Accession: AC2796
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A;Cross-references: UNIPROT:Q8UEH1; UNIPARC:UPI00001643A5;
C;Genetics:
A;Gene: AGR_C_3284
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C; Superfamily: 9
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Ferrandez A., Minambres B., Garcia B., Olivera E.R., Luengo J.M.

Garcia J.L., Diaz E.;

"Catabolism of phenylacetic acid in Escherichia coli. Character:

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27-APR-2001, integrated into UniProtKB/Swiss-Prot.
01-FEB-1997, sequence version 1.
07-MAR-2006, entry version 56.
Probable phenylacetic acid degradation NADH oxidor
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     NUCLEOTIDE SEQUENCE
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nitted (APR-2004) to the EMBL/GenBank/DDBJ databases.
CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
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PROSITE; PR00197; 2FE2S_FER_1; 1.

PROSITE; PS01097; 2FE2S_FER_2; 1.

PROSITE; PS51085; 2FE2S_FER_2; 1.

2FE-2S; Complete proteome; Electron transport; FAD; Flavoprotein; 2FE-2S; Complete proteome; Electron transport; FAD; Pransport.

Probable phenylacetic acid degradatic phenylacetic acid degradatic NADH oxidoreductase paaE.
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InterPro; IPR001041; Ferredoxin.
InterPro; IPR012675; Ferredoxin fold.
InterPro; IPR012675; Ferredoxin fold.
InterPro; IPR001709; FPN cyt_redctse.
InterPro; IPR008333; Oxred FAD bd.
InterPro; IPR001433; Oxred FAD bd.
InterPro; IPR001433; Oxred FAD bd.
InterPro; IPR011884; PA CA Oase5.
InterPro; IPR011221; Phe hydroxylase.
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LinkHub; P76081; -.
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EMBL; U00096; AAC74474.1; -; Genomic_DNA.
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EMBL; D90778; BAA15003.1; AUT_INIT; Genom.
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MEDLINE=20229831; PubMed=10766858; DOI=10.1074/jbc.275.16.12214;
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Science 277:1453-1474(1997).
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Nucleic Acids Res. 28:2311-2314(2000).
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01-MAY-1999, sequence version 1.
01-MAY-1999, sequence version 1.
07-MAR-2006, entry version 36.
UDP-N-acetylglucosamine--N-acetylmuramyl-(pentapeptide)
UPP-N-acetylglucosamine-N-acetylglucosamine transferase
pyrophosphoryl-undecaprenol N-acetylglucosamine transferase
(EC 2.4.1.227) (Undecaprenyl-PP-MurNAc-pentapeptide-UDPGlcNAc (
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                                                           NUCLEOTIDE SEQUENCE
STRAIN=TW-183;
Geng M.M., Schuhmach
                                                                                                                                                                                                                                                                                         MEDLINE=20150255; PubMed=10684935; DOI=10.1093/nar/28.6.1397; Read T.D., Brunham R.C., Shen C., Gill S.R., Heidelberg J.F., White O., Hickey E.K., Peterson J.D., Utterback T.R., Berry K.J. Bass S., Linher K.D., Weidman J.F., Khouri H.M., Craven B., Bown Dodson R.J., Gwinn M.L., Nelson W.C., DeBoy R.T., Kolonay J.F., McClarty G., Salzberg S.L., Eisen J.A., Fraser C.M., Salzberg S.L., Eisen J.A., Fraser C.M.
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Geng M.M., Schuhmacher A., Muehldorfer I., Bensch K.W., Schaefer K.P., Schneider S., Pohl T., Essig A., Marre R., Melchers K.; The genome sequence of Chlamydia pneumoniae TW183 and comparison with other Chlamydia strains based on whole genome sequence analysis."; Submitted (MAY-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                        MEDLINE=20330349; PubMed=10871362; DOI=10.1093/nar/28.12.2311;
Shirai M., Hirakawa H., Kimoto M., Tabuchi M., Kishi F., Ouchi
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                                                                                                                                                                                                                                                   "Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.";
Nucleic Acids Res. 28:1397-1406(2000).
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                                                                                                                                 Q5SMX2;
21-DEC-2004, integrated into Uni
21-DEC-2004, sequence version 1
21-DEC-2006, entry version 7.
07-FEB-2006, entry version 7.
Basic helix-loop-helix protein-
Name=90498801.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TIGR; CP0962; -...
BioCyc; CPNE115711:CP0962-MONOMER; -..
BioCyc; CPNE115713:CPN0904-MONOMER; -..
BioCyc; CPNE138677:CPJ0904-MONOMER; -..
BioCyc; CPNE138677:CPJ0904-MONOMER; -..
BioCyc; CPNE182082:CPB0936-MONOMER; -..
HAMAP; MF 00033; -; 1.
InterPro; IPR007235; Glyco_tran_28_C.
InterPro; IPR004276; Glyco_trans_28_C.
InterPro; IPR004276; Murg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AE001670; AAD19042.1; -; Genomic DNA.
EMBL; AE002254; AAF38742.1; -; Genomic DNA.
EMBL; BA000008; BAA99112.1; -; Genomic DNA.
EMBL; AE017160; AAP98865.1; ALT_INIT; Genomic DNA.
EMBL; C72022; C72022.
PIR; F86603; F86603.
                                                                                                                                                                                                                                                                                  X2 ORYSA
Q5SMX2 ORYSA
                                              Oryza sativa (japonica cultivar-group).
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; BEP cla
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF04101; Glyco_tran_28_C; 1.
Pfam; PF03033; Glyco_transf 28; 1.
TIGRFAMs; TIGR01133; murg; I.
Cell cycle; Cell division; Cell shape; Cell wa
Glycosyltransferase; Inner membrane; Membrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
  NCBI_TaxID=39947; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GenomeReviews;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GenomeReviews;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptidoglycan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         subunit on undecaprenyl-pyrophosphoryl-murNAc-pentapeptide (lipid intermediate I) to form undecaprenyl-pyrophosphoryl-murNAc-pentapeptide (lipid intermediate II) to form undecaprenyl-pyrophosphoryl-MurNAc-(pentapeptide)GlCNAc (lipid intermediate II) (By similarity).

- CATALYTIC ACTIVITY: UDP-N-acetylglucosamine + MurZAc (cyyl-L-Ala-gamma-D-Glu-L-Lys-D-Ala-D-Ala)-diphosphoundecaprenol = UDP + GlCNAc-(l-x4)-MurZAc(cyyl-L-Ala-gamma-D-Glu-L-Lys-D-Ala)-diphosphoundecaprenol.

- PATHWAY: Peptidoglycan biosynthesis; last step.

SUBCELLULAR LOCATION: Bacterial cell inner membrane; peripheral membrane protein (By similarity)
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; AE001363 GR;
; BA000008 GR;
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1 357
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CpB0936.
CPn0904.
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Pred. No. 91;
Mismatches
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undecaprenol N-acetylglucosamine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UDP-N-acetylglucosamine--N-acetylmuramyl-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /FTId=PRO
                                                                                                                                                                                                                                                                                                    PRT;
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                                                                             BEP clade;
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RESULT 6
Q98L45 RHILO
ID 098L45;
pT 01-0CT-2001, s
pT 01-0CT-2001, s
pT 01-CCT-2001, s
pT 01-CCT-2001,
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RA Wu J., Niimura Y., Cheng Z., Nagamura Y., Antonio B.A., Kanamori H.,
RA Hosokawa S., Masukawa M., Arikawa K., Chiden Y., Hayashi M.,
RA Okamoto M., Ando T., Aoki H., Arita K., Hamada M., Harada C.,
RA Hijishita S., Honda M., Ichikawa Y., Idonuma A., Iijima M., Ikeda M.,
RA Ikeno M., Ito S., Ito T., Ito Y., Ito Y., Iwabuchi A., Kamiya K.,
RA Karasawa W., Katagiri S., Kikuta A., Kobayashi N., Kono I.,
RA Nagasaki H., Nakashima M., Nakama Y., Nakamichi Y., Nakamura M.,
RA Nagasaki H., Nakashima M., Nakama Y., Nakamichi Y., Nakamura M.,
RA Namiki N., Negishi M., Ohta I., Ono N., Saji S., Sakai K., Shibata M.,
RA Shimokawa T., Shomura A., Song J., Takazaki Y., Terasawa K., Tsuji K.,
A Waki K., Yamagata H., Yamane H., Yoshiki S., Yoshihara R., Yukawa K.,
A Yano M., Jiang J., Gojobori T.;
Ti "The Genome accusero. Sai T. T.,
Ti The Genome accusero. Sai T. T.,
Ti Terasawa K., Tsuji Y.,
Ti Terasawa H., Endo T., Ito H., Hahn J.H., Kim H.-I., Eun M.-Y.,
Ti The Genome accusero. Sai T. T.,
Ti The Genome accusero. Sai T. T.,
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Ti Terasawa K., Tsuji Y.,
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Best Local
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GO; GO:0005634; F:transcription regulator activity;
GO; GO:0045449; F:regulation of transcription; IEA.
InterPro; IPR001092; HLH basic.
InterPro; IPR011598; HLH_DNA_bd.
Pfam; PF00010; HLH; 1.
SMART; SM00353; HLH; 1.
PROSITE; PS50888; HLH; 1.
PROSITE; PS50888; HLH; 1.
                                                                                                                                                                                                                                                  "Complete genome structure of the nitrogen-fixing Mesorhizobium loti.";
                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=21082930; PubMed=11214968; DOI=10.1093/dnares/7.6.331; Kaneko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasamoto Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T., Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A., Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M., Takeuchi C., Yamada M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OrderedLocusNames=m111183;
Rhizobium loti (Mesorhizobium loti).
Ratiobium loti (Mesorhizobium loti).
Bacteria; Proteobacteria; Alphagroteobacteria; Rhizobiales;
Phyllobacteriaceae; Mesorhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE STRAIN=MAFF303099;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2001, sequence version 21-FEB-2006, entry version 22
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Nature 420:312-316(2002).
-!- SIMILARITY: Contains 1 basic helix-loop-helix (bHLH) domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE
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                                                     by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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Pred. No.
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Best Local S
Matches 6
Query Match
Best Local Similarity
                                                                                                       InterPro; IPR003594; ATP_bd_ATPASE.
InterPro; IPR00458; His_kin_like_C.
InterPro; IPR003661; His_kin_A.
InterPro; IPR003667; His_kin_Be.
InterPro; IPR003667; His_kinase.
InterPro; IPR00348; Lantibiot_regn.
Pfam; PP00518; HATPASE c; 1.
Pfam; PF00512; HiSKA; T.
Pfam; PF00512; HiSKA; T.
PF10736; HATPASE c; 1.
PR10736; LANITAEGEPROT.
SMART; SM00387; HATPASE c; 1.
SMART; SM00387; HATPASE c; 1.
SMART; SM00388; HISKA; T.
SMOSITE; PS50109; HIS KIN; 1.
SMART; SM00388; HISKA; 1.
SMOSITE; PS50109; HIS KIN; 1.
SEQUENCE 432 AA; 48532 MW; 74A8E90.
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PROSITE; PS50975; ATP GRASP; 1. Nucleotide binding; INTERPRO; IPR011761; ATP GRASP; INTERPROSITE; PS50975; ATP GRASP; 1. Nucleotide binding; INTERPROSITE; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0000155; F:two-component sensor molecule activity; IEA. GO; GO:0006468; P:protein amino acid phosphorylation; IEA. GO; GO:0000160; P:two-component signal transduction system (p.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AY548383; AAT12782.1; -; Genomic_
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0016301; F:kinase activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=15640200; DOI=10.1128/AEM.71.1.290-296.2005; Wilson M.M., Metcalf W.W.; Wilson M.M., Metcalf W.W.; Oxidation of Reduced Phosphorus Compounds by Alcaligenes faecal:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-FEB-2005, integrated into UniProtKB/TrEMBL.
15-FEB-2005, sequence version 1.
07-FEB-2006, entry version 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/t.Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wilson M., Metcalf W.;
Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-WM20
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Q5J1L5;
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6; Conserv
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46722 MW; F4435AC1EDC7F645 CRC64;
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Pred. No. 1.
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Pred. No.
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                                   Length 432;
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Q45AY3 9BURK
Q45AY3;
Q45AY3;
13-SEP-2005, integrated into UniProtKB/T
13-SEP-2005, sequence version 1.
13-SEP-2006, entry version 5.
Q7-FEB-2006, entry version 5.
Amino acid permease-associated region.
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Q62CX5 F
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005279; F:amino acid-polyamine transporter
GO; GO:000665; P:amino acid transport; IEA.
GO; GO:000665; P:transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR002293; AA/rel_permease1.
InterPro; IPR004841; Permease_region.
PANTHER; PTHR1785; AA/rel_permease1; 1.
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PubMed=15377793; [
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Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
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25-OCT-2004,
                                                                              ORFNames=BcenDRAFT 5191;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00324; AA_permease; 1.
PROSITE; PS00218; AMINO_ACID_PERMEASE_1; 1.
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                                     Burkholderia cenocepacia AU 1054.
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447 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative 0
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                                                                                                                                                                                                                       integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAU46888.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      σ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 48683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MW;
   Betaproteobacteria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A209ED83F6C1D427 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                          460
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . 1.2e+02;
ches 0;
                                                                                                                                                                                                                                                                                          A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽
   Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 447;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT QALTZ22 ID TQALTZ22 ID TQALTZ22 ID TQALTZ22 DT 022 
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Z2_9BURK
Q4LTZ2_9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=AU 1054;
US DOE Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
"Sequencing of the draft genome assembly of E
AU 1054.";
NUCLEOTIDE SEQUENCE.
STRAIN=HI2424;
US DOE Joint Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00324; AA permease; 1.

Pfam; PF00324; AA permease; 1.

PROSITE; PS00218; AMINO ACID PERMEASE 1; 1.

PROSITE; PS00218; AMINO ACID PERMEASE 1; 1.

SEQUENCE 460 AA; 49648 MW; 68780B9E9F03369D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                  "Sequencing HI2424.";
                                                                                                                                                                                                            Copeland A., Lucas S., Lapidus A., Barry K., Hammon N., Israni S., Pitluck S., Richardson "Sequencing of the draft genome assembly of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales; Burkholderiaceae; Burkholderia; Burkholderia cepacia complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Burkholderia cenocepacia HI2424.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amino acid permease-associated region. ORFNames=Bcen2424DRAFT_4131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-AUG-2005, sequence version 07-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-AUG-2005, integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q4LTZ2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AU 1054.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Larimer F., Land M.; "Annotation of the draft
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Burkholderiaceae; Burkholderia; Burkholderia cepacia complex
NCBI_TaxID=331271;
                                                                                                                                                 Submitted
                                                                                                                                                                                                                                                                                                                        US DOE Joint Genome Institute (JGI-PGF);
                                                                                                                                                                                                                                                                                                                                                                 STRAIN=HI2424;
                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=331272;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR002293; AA/rel_permease1
InterPro; IPR004840; AAc_permease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US DOE Joint Genome Institute (JGI-ORNL);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-AU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO:0016021; C:integral to membrane;
GO:0016020; C:membrane; IEA.
GO:0005279; F:amino acid transport;
GO:0006865; P:amino acid transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO:0006810;
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       Institute
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       (JGI-ORNL);
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; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IEĀ.
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P.;
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RESULT 1

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GO; GO:0016020; C:membrane; IEA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005279; F:amino acid-polyamine transporter activity; IEA.
GO; GO:0006865; P:amino acid transport; IEA.
GO; GO:0006810; P:transport; IEA.
R InterPro; IPR002293; AA/rel_permease1.
R InterPro; IPR002293; AA/rel_permease.
R InterPro; IPR002293; AA/rel_permease.
R InterPro; IPR002293; AA/rel_permease.
R PANTHER; PTHR11785; AA/rel_permease1; 1.
DR PF00324; AA permease; 1.
DR PF00324; AA permease; 1.
DR PROSITE; PS00218; AMINO_ACID_PERMEASE 1; 1.
DR PROSITE; PS00218; AMINO_ACID_PERMEASE 1; 1.
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                                                                Query Match
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Q395V1_BURS3
Q395V1;
22-NOV-2005,
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                                                                                                                                                                                                            88888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US DOE Joint Genome Institute;
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina T.,
Hammon N., Israni S., Pitluck S., Chain P., Malfatti S., Shin M.,
Vergez L., Schmutz J., Larimer F., Land M., Kyrpides N., Lykidis A.,
Richardson P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amino acid transporter.
OrderedLocusNames=Bcep18194_B1646;
Burkholderia sp. (strain 383) (Bur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (JU-!- CAUTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Complete sequence of chromosome 2 of Burkhold Submitted (OCT-2005) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
Burkholderiaceae; Burkholderia; Burkholderia cepacia complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-NOV-2005, sequence version 21-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                  EMBL;
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                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCIB 9086 / R18194)).
Local Similarity
les 6; Conserv
                                                                                                                                                                                                         L; CP000152; ABB11760.1; -; Genomic_DNA.
GO:0016021; C:integral to membrane; IEA.
GO:0016020; C:membrane; IEA.
GO:0005279; F:amino acid-polyamine transporter activity; IEA.
GO:0006865; P:amino acid transport; IEA.
GO:0006810; P:transport; IEA.
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EMBL/GenBank/DDBJ whole
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460 AA;
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of the d
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   Conservative
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                                  Score 31; DB 2;
Pred. No. 1.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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                                                                                                                                            D6BBAA9AF2CD5155
   Mismatches
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1.2e+02;
thes 0;
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R3 BURP1
Q3JFR3 BURP1
Q3JFR3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fraser C.M., Casjens S., Huang W.M., Sutton G.G., Clayton R.A., Lathigra R., White O., Ketchum K.A., Palmer N., Dodson R., Liathigra R.K., Gwinn M., Dougherry B., Fleischmann R.D., Richardson D., Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M., van-Vugt R., Adams M.D., Gocayne J.D., Weidman J., Utterback T., Watthey L., McDonald L., Artiach P., Bowman C., Garland S., Fujii C., Cotton M.D., Horst K., Tomb J.-F., Roberts K., Hatch B., Smith H.O., Venter J.C.;
                                                                                                                                           TIGR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amino acid permease.
ORFNames=BTH_II1560;
Burkholderia thailandensis
                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
Woods D.E., Nierman W.C.;
Submitted (SEP-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-NOV-2005, integrated into UniProtKB/TrEMBL 08-NOV-2005, sequence version 1. 21-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=320372;
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Burkholderiaceae; Burkholderia; pseudomallei group.
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Q2T4Z4;
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BURTH
L; CP000125; ABA52590.1; -; Genomic_DNA.
R; BURPSI710b_A2440; -.
GO:0016021; C:integral to membrane; IEA.
GO:0016020; C:membrane; IEA.
GO:0005279; F:amino acid-polyamine transporter
GO:0006865; P:amino acid transport; IEA.
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VCE 461 AA; 49957 MW; 7F1195E772F28B65
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                                         activity; IEA
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Best Local
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InterPro; IPR002293; AA/rel permeasel.
InterPro; IPR004840; AAC.permease.
InterPro; IPR004841; Permease region.
Pfam; PF00324; AA permease; 1.
PROSITE; PS00218; AMINO_ACID_PERMEASE_1;
Complete proteome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=15377794; DOI=10.1073/pnas.0403302101;
Holden M.T.G., Titball R.W., Peacock S.J., Cerdeno-Tarraga A.-M.,
Atkins T., Crossman L.C., Pitt T., Churcher C., Mungall K.L.,
Bentley S.D., Sebaihia M., Thomson N.R., Bason N., Beacham I.R.,
Brooks K., Brown K.A., Brown N.F., Challis G.L., Cherevach I.,
Chillingworth T., Cronin A., Crossett B., Davis P., DeShazer D.,
Feltwell T., Fraser A., Hance Z., Hauser H., Holroyd S., Jagels K.,
Keith K.E., Maddison M., Moule S., Price C., Quail M.A.,
Keith K.E., Maddison M., Moule S., Price C., Quail M.A.,
Kentheriotic E., Rutherford K., Sanders M., Simmonds M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name=proY; OrderedLocusNames=BPSS0845;
Burkholderia pseudomallei (Pseudomonas pseudomallei).
Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-OCT-2004, integrated into Unil
25-OCT-2004, sequence version 1.
07-FEB-2006, entry version 17.
                                                                                                                                                                                                          InterPro; IPR004841; Permease_region. PANTHER; PTHR11785; AA/rel_permease1; 1.
                                                                                                                                                                                                                                                 GG; GG:0016021; C:integral to membrane; IEA.
GG; GG:0016020; C:membrane; IEA.
GG; GG:0005279; F:amino acid-polyamine transporter activity; IEA.
GG; GG:000585; P:amino acid transport; IEA.
GG; GG:0006810; P:transport; IEA.
InterPro; IPR00293; AA/rel permease1.
InterPro; IPR004840; AAc_permease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Burkholderia pseudomallei."; Proc. Natl. Acad. Sci. U.S.A. 101:14240-14245(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=K96243
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PROSITE; PS00218; AMINO_ACID_PERMEASE_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Distributed under the Creative Commons Attribution-NoDerivs License
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BURPS
                                                                               Local
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                                                        Similarity 6; Conserv
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Pred. No. 1.2e+02;
                                                                           Score 31; DB 2;
Pred. No. 1.2e+02
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Azevedo V., F
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Submitted (JUN-1997)
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"The complete genome
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RA Azevedo V. Bertero M.G. Bessieres P. Bolotini A. Borchert S.,
RA Borriss R., Boursier L. Brans A., Braun M., Brignell S.C., Bron S.,
RA Borriss R., Boursier L. Brans A., Braun M., Brignell S.C., Bron S.,
RA Borriss R., Boursier L., Brans A., Braun M., Brignell S.C., Bron S.,
RA Borriss R., Boursier L., Brans A., Braun M., Brignell S.C., Bron S.,
RA Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
RA Choi S.-K., Codani J.-J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.-K., Devine K.M., Duesterhoeft A., Ehrlich S.D., Emmerson P.T.,
RA Britz C., Fujita M., Fujita Y., Fanger C., Ferrari E., Foulger D.,
RA Ghim S.-Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Ghim S.-Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Gilm S.-Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Hilbert H., Holsappel S., Hosono S., Hullo M.-F., Itaya M.,
RA Jones L.-M., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M.,
RA Jones L.-M., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M.,
RA Medigue C., Kobayashi Y., Koetter P., Koningstein G., Krogh S.,
RA Kumano M., Kurita K., Lapidus A., Liu H., Masuda S., Mauel C.,
RA Kumano M., Kurita K., Lapidus A., Liu H., Masuda S., Mauel C.,
RA Medigue C., Medina N., Mellado R.P., Liu H., Masuda S., Mauel C.,
RA Medigue C., Medina N., Mellado R.P., Miruno M., Moestl D., Nakai S.,
RA Park S.-H., Parro V., Pohl T.M., Portetelle D., Porwollik S.,
RA Park S.-H., Parro V., Pohl T.M., Portetelle D., Porwollik S.,
RA Rey M., Reynolds S., Rieger M., Rivolta C., Rocha E., Roche B.,
RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R.,
RA Rose M., Sadaie V., Sato T., Scanlan E., Schleich S., Schroeter R.,
RA Rose M., Vasainchi M., Tamakoshi A., Tanaka T., Terpstra P.,
RA Wassarotti A., Wahuters P., Wipat A., Yamamoto H., Yamane K.,
RA Yashikawa H., Danchin A., Landelle E., Roche E.,
RA Yashikawa H., Danchin A., Landelle E., Koshikawa H.-F., Zumstein E.,
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O34618; Q795E5;
O1_JAN_1998, integrated into UniProtKB/TrEMBL.
O1_JAN_1998, sequence version 1.
O7_FEB-2006, entry version 30.
Amino acid transporter (YthA protein).
Name=ythA; OrderedLocusNames=BSU30530;
Bacillus subtilis.
Paranceia. Simminose Bacillales Bacillaceae
EMBL; Z99119; CAB15031.1; -; Genomic_DNp
PIR; C69997; C69997.
BioCyc; BSUB1423:BSU3048-MONOMER; -.
GO; GO:0016021; C:integral to membrane;
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005279; F:amino acid-polyamine t
                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
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MEDLINE=98044033; PubMed=9384377; DOI=10.1038/36786;
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Yoshikawa H., Danchin A.;
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MEDLINE=98048467; PubMed=9387221;
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                                                                                                                                                                                                                                               AF008220; AAC00244.1; -;
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Bessieres P., Bolotin A., Borche
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n of the
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                                                                                        IEA.
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RESULT 16
Q2STX5 BURN
Q2STX5 DIR
AC Q2STX
AC Q2S
RESULT 17
Q3JY99 BURP1
ID Q3JY99;
AC Q3JY99;
DT 08-NOV-2005, s
DT 21-FEB-2006, c
DE Amino acid per
GN OrderediocusNa
OS Burkholderia proto
OC Burkholderiace
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Best Local S
Matches 6
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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Lathigra R., White O., Ketchum K.A., Palmer N., Dodson R.,
Hickey E.K., Gwinn M., Dougherty B., Fleischmann R.D., Richardson D.,
Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M.,
van-Vugt R., Adams M.D., Gocayne J.D., Weidman J., Utterback T.,
Watthey L., McDonald L., Artiach P., Bowman C., Garland S., Fujil C.,
Cotton M.D., Horst K., Tomb J.-F., Roberts K., Hatch B., Smith H.O.,
Venter J.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GO; GO:0006865; P:amino acid transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR002293; AA/rel_permease1.
InterPro; IPR004840; AAc_permease.
InterPro; IPR004841; Permease_region.
PANTHER; PTHR11785; AA/rel_permease1; 1.
Pfam; PF00324; APPERMEASE; 1; UNKNOWN_1.
Complete proteome; Transmembrane; Transport.
SEQUENCE 463 AA; 50328 MW; B21F9F40AB1C7180 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amino acid permease.
ORFNames=BTH_I3129;
Burkholderia thailandensis
                                                                                Amino acid permease.
OrderedLocusNames=BURPS1710b_0036;
                                                                                                                                08-NOV-2005, integrated into UniProtKB/TrEMBL. 08-NOV-2005, sequence version 1. 21-FEB-2006, entry version 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JAN-2006, sequence version 07-FEB-2006, entry version 2.
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Q2STX5;
                           Burkholderia pseudomallei (strain 1710b).
Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (JUL-2005) to the EMBL/GenBank/DDBJ databases.
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NCE 468 AA; 50779 MW;
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Pred. No. 1.2
); Mismatches
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Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genomic_DNA.
A905AA5CB57B7284 CRC64;
     pseudomallei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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R GO; GO:0016021; C:integral to membrane; IEA.

R GO; GO:0016020; C:membrane; IEA.

R GO; GO:0005279; F:amino acid-polyamine transporter activity; IEA.

R GO; GO:0006865; P:amino acid transport; IEA.

R GO; GO:0006810; P:transport; IEA.

R GO; GO:0006810; P:transport; IEA.

R InterPro; IPR002293; AA/rel permease1.

R InterPro; IPR004840; AAc_permease.

R InterPro; IPR004841; Permease_region.

R Pfam; PF00324; AA permease 1.

R PROSITE; PS00218; AMINO_ACID_PERMEASE_1; 1.

R PROSITE; PS00218; AMINO_ACID_PERMEASE_1; 1.
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Best Local (
TIGR; BMAJ796; -.

GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005020; C:membrane; IEA.
GO; GO:0005279; F:amino acid transport; IEA.
GO; GO:0006865; P:amino acid transport; IEA.
GO; GO:0006810; P:transport; Dermease1.

InterPro; IPR002293; AA/rel_permease.

InterPro; IPR003840; AAc_permease.

InterPro; IPR003840; AAc_permease.
                                                                                                                                                                                                                                                                                                                                                                                            Nierman W.C., DeShazer D., Kim H.S., Tettelin H., Nelson K.E., Reldblyum T.V., Ulrich R.L., Ronning C.M., Brinkac L.M., Daugherty S.C., Davidsen T.D., DeBoy R.T., Dimitrov G., Dodson Durkin A.S., Gwinn M.L., Haft D.H., Khouri H.M., Kolonay J.F., Madupu R., Mohammoud Y., Nelson W.C., Radune D., Romero C.M., Sarria S., Selengut J., Shamblin C., Sullivan S.A., White O., Y Zafar N., Zhou L., Fraser C.M.; "Structural flexibility in the Burkholderia mallei genome."; "Structural flexibility in the Burkholderia mallei genome.";
                                                                                                                                                                                                                                                               EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burkholderia mallei (Pseudomonas mallei).
Bacteria; Proteobacteria; Betaproteobacte:
Burkholderiaceae; Burkholderia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-OCT-2004, integrated into 0
25-OCT-2004, sequence version
07-FEB-2006, entry version 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed=15377793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OrderedLocusNames=BMA2796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amino acid permease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q62G62;
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                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [LARGE SCALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=13373;
                                                                                                                                                                                                                                                                                                                  Distributed under the Creative Commons Attribution-NoDerivs
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BURMA
                                                                                                                                                                                                                                                                  CP000010; AAU47957.1; -;
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Pred. No. 1.2
D; Mismatches
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                                                                                                                                                                                                                                                               Genomic_DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 468;
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                                                                                                                                                            activity;
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RESULT 19
Q63PW6 BURPS
Q63PW6 BURPS
AC Q63PW6;
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Matches 6
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Holden M.T.G., Titball R.W., Peacock S.J., Cerdeno-Tarraga A.-M., Atkins T., Crossman L.C., Pitt T., Churcher C., Mungall K.L., Bentley S.D., Sebaihia M., Thomson N.R., Bason N., Beacham I.R., Brooks K., Brown K.A., Brown N.F., Challis G.L., Cherevach I., Chillingworth T., Cronin A., Crossett B., Davis P., DeShazer D., Feltwell T., Fraser A., Hance Z., Hauser H., Holroyd S., Jagels K., Keith K.E., Maddison M., Moule S., Price C., Quail M.A., Keith K.E., Maddison M., Moule S., Sanders M., Samonds M., Songsivilai S., Stevens K., Tumapa S., Vesaratchavest M., Shitchead S., Yeats C., Barrell B.G., Oyston P.C.F., Parkhill J., "Genomic plasticity of the causative agent of melioidosis, Burkholderia pseudomallei":
                                                                                                                                                                                                                                                                                                                                         EMBL; BX571965; CAH37269.1; -; Genomic_DNA.
GO; GO:0016021; C:integral to membrane; IEA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:00165279; F:amino acid-polyamine transporter activity; IEA.
GO; GO:0005879; F:amino acid transport; IEA.
GO; GO:0006865; P:amino acid transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR004840; AA/rel_permease1.
InterPro; IPR004840; AA/rel_permease.
InterPro; IPR004841; Permease_region.
PANTHER; PTHA1785; AA/rel_permease1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        063PW6;
25-OCT-2004, integrated into Unil
25-OCT-2004, sequence version 1.
07-FEB-2006, entry version 16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PANTHER; PTHR11785; AA/rel_permease1; 1.
Pfam; PF00324; AA permease; 1.
PROSITE; PS00218; AMINO_ACID_PERMEASE_1;
Complete protecome.
SEQUENCE 468 AA; 50949 MW; AAEEB96964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Holden M.T.G., Titball R.W., Peacoc
Atkins T., Crossman L.C., Pitt T.,
                                                                                                                                                                                                                                                                                     Pfam; PF00324; AA_permease; 1.
PROSITE; PS00218; AMINO_ACID_PERMEASE_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Burkholderia pseudomallei.";
Proc. Natl. Acad. Sci. U.S.A. 101:14240-14245(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=15377794; DOI=10.1073/pnas.0403302101; Holden M.T.G., Titball R.W., Peacock S.J., Ce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=K96243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales; Burkholderiaceae; Burkholderia; pseudomallei group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Burkholderia pseudomallei (Pseudomonas pseudomallei)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OrderedLocusNames=BPSL3256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Putative amino acid permease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=28450;
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                                                                                                               Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FSRYAR 6
                                                         FSRYAR
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   FSRYAR
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                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             integrated into UniProtKB/TrEMBL.
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Pred. No. 1.2
); Mismatches
                                                                                                                                       Score 31; DB 2; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
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                                                                                                                                                                                                                                  69E6B0D62E681105 CRC64;
                                                                                                                  Mismatches
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                                                                                                                                                                      Length 468;
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RESULT 20

RESCULT ID ASCB EACOLI

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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1997, sequence version 2. 07-MAR-2006, entry version 51. 6-phospho-beta-glucosidase ascB (EC Name=ascB; OrderedLocusNames=b2716;
                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS00572; GLYCOSYL_HYDROL_F1_1; 1.
PROSITE; PS00653; GLYCOSYL_HYDROL_F1_2; 1.
Complete proteome; Glycosidase; Hydrolase.
                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001360; Glyco hydro 1.
PANTHER; PTHR10353; Glyco hydro 1; 1.
Pfam; PF00232; Glyco hydro 1; 1.
PRINTS; PR00131; GLHYDRLASE1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EcoGene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HSSP; P11546; 1PBG.
GenomeReviews; U00096_GR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; M73326; AAA16430.1; -; Unassigned_DNA.
EMBL; U09579; AAA69226.1; ALT INIT; Genomic_DNA.
EMBL; U0096; AAC75758.1; -; Genomic_DNA.
PIR; H65051; H65051.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Science
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mau B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=97426617; PubMed=9278503; DOI=10.1126/science.277.5331.1453; Blattner F.R., Plunkett G. III, Bloch C.A., Perna N.T., Burland V., Riley M., Collado-Viddes J., Glasner J.D., Rode C.K., Mayhew G.F., Gregor J., Davis N.W., Kirkpatrick H.A., Goeden M.A., Rose D.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE
STRAIN=K12
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01-MAR 1992, integrated into UniProtKB/Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BioCyc; EcoCyc:EG10085-MONOMER;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EchoBASE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H(2)0 = D-glucose + D-glucose 6-phosphate.
-i- SIMILARITY: Belongs to the glycosyl hydrolase 1 family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- FUNCTION: Can hydrolyze salicin, cellobiose, and probably arbutin.
-!- CATALYTIC ACTIVITY: 6-phospho-beta-D-glucosyl-(1,4)-D-glucose +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The complete genome sequence of Escherichia coli K-12.";
Science 277:1453-1474(1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hall B.G., Xu L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=92334140; PubMed=1630307;
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                                                         6; Conserv
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EG10085; ascB.
  FSRYAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                              Conservative
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                                                                                                                                                                                                                                 6-phospho-beta-glucosidase ascB.
(FTId=PRO 000063895.
Proton donor (Potential).
Nucleophile (By similarity).
EA -> GT (in Ref. 1).
                                                                                                                                                                                     EA -> GT (in Ref. 1)
S -> C (in Ref. 1).
RK -> HR (in Ref. 1
                                                                                   Score 31; I
Pred. No. 1
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                                                                                                                                                              ( -> HR (in Ref. 1)
02ACE6BEBF211011 C
                                                                Mismatches
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FSRYAR 162

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RESULT 22
Q32CL4_SHIDS
ID Q33CL4_SHIDS
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PubMed=16275786; DOI=10.1093/nar/gki954;
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Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
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                                                                                                                                                                                                                      Yang F., Yang J., Zhang X., Chen L., Jiang Y., Yan Y., Wang J., Xiong Z., Dong J., Xue Y., Zhu Y., Xu X., Sun Nie H., Peng J., Xu J., Wang Y., Yuan Z., Wen Y., Yao Qiang B., Hou Y., Yu J., Jin Q., "Genome dynamics and diversity of Shigella species, the agents of bacillary dysentery.";

Nucleic Acids Res. 33:6445-6458(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=16275786; DOI=10.1093/nar/gki954;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6-phospho-beta-glucosidase.
Namme-ascB; OrderedLocusNames=SDY 2912;
Shigella dysenteriae serotype 1 (strain Sd197).
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-DEC-2005, integrated into UniProtKB/TrEMBL 06-DEC-2005, sequence version 1. 07-FEB-2006, entry version 4.
                                                                                                                                       Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                GO; GO:0004553;
GO; GO:0005975;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=300267;
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GO:0004553; F:hydrolase activity, hydrolyzing O-glycosyl
GO:0005975; P:carbohydrate metabolism; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 157
                                                          CP000034; ABB62941.1; -; Genomic_DNA.
O:0004553; F:hydrolase activity, hydrolyzing O-glycosyl
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                                P:carbohydrate
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Pred. No.
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                                   metabolism;
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L., Chen S.,
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E8 SHISS
PRELIMINARY; PRT; 474 AA.

Q3YYE8;
Q3YYE8;
27-SEP-2005, integrated into UniProtKB/TrEMBL.
27-SEP-2005, sequence version 1.
07-FEB-2006, entry version 5.
07-FEB-2006, entry version 5.
Name-ascB; OrderedlocusNames=SSO_2860; ORFNames=SSO_2860;
Shigella sonnei (strain Ss046).
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

PubMed=16275786; DOI=10.1093/nar/gki954;

Yang F., Yang J., Zhang X., Chen L., Jiang Y., Yan Y., Tang X.,

Wang J., Xiong Z., Dong J., Xue Y., Zhu Y., Xu X., Sun L., Chen S.

Nie H., Peng J., Xu J., Wang Y., Yuan Z., Wen Y., Yao Z., Shen Y.,

Qiang B., Hou Y., Yu J., Jin Q.;

"Genome dynamics and diversity of Shigella species, the etiologic agents of bacillary dysentery.";

Nucleic Acids Res. 33:6445-6458(2005).
                                                                                                                                                                   Q2MAB3;
21-FEB-2006, integrated into UniProtKB/TrEMBL
21-FEB-2006, sequence version 1.
07-MAR-2006, entry version 2.
Cryptic 6-phospho-beta-glucosidase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; CP000038; AAZ89464.1; -; Genomic_DNA.
GO; GO:0004553; F:hydrolase activity, hydrolyzing O-glycosyl
GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR001360; Glyco_hydro_1.
Pfam; PF00332; Glyco_hydro_1; 1.
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PROSITE; PS00572; GLYCOSYL_HYDROL_F1_1;

PROSITE; PS00653; GLYCOSYL_HYDROL_F1_2;
                                          NCBI_TaxID=316407;
                                                                                                 Bacteria; Proteobacteria;
                                                                                                                         Escherichia coli W3110.
                                                                                                                                                      Name=ascB
                                                                                                                                                                                                                                                                                                    QZMAB3_ECOLI
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                                                                         Enterobacteriaceae;
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  NUCLEOTIDE SEQUENCE
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ECOLI
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Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
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Attribution-NoDerivs License
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MEDINE-97349980; PubMed-9205837; DOI=10.1093/dnares/4.2.91; MEDINE-97349980; PubMed-9205837; DOI=10.1093/dnares/4.2.91; Yamamoto Y., Aiba H., Baba T., Hayashi K., Inada T., Isono K., Itoh T., Kimura S., Kitagawa M., Makino K., Miki T., Mitsuhashi N. Mizobuchi K., Mori H., Nakade S., Nakamura Y., Nashimoto H., Oshima T., Oyama S., Saito N., Sampei G., Satoh Y., Sivasundaram Tagami H., Takahashi H., Takeda J., Takemoto K., Uehara K., Wada "Construction of a contiguous 874-kb sequence of the Escherichia -Kl2 genome corresponding to 50.0-68.8 min on the linkage map and analysis of its sequence features.";
                                                                                                                                                                                                                                                                                                                                    PubMed=16397293; DOI=10.1093/nar/gkj150; Riley M., Abe T., Arnaud M.B., Berlyn M.K., Blattner F.R Chaudhuri R.R., Glasner J.D., Horitari T., Keseler I.M., Mori H., Perna N.T., Plunkett G. III, Rudd K.E., Serres Thomas G.H., Thomson N.R., Wishart D., Wanner B.L.; "Escherichia coli K-12: a cooperatively developed annota
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Ohtsubo E., Baba T., Wan
Ohtsubo accurate genome
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                                                                                                                                     -KI2 genome corresponding analysis of its sequence f DNA Res. 4:91-113/1007
                                                                                                                                                                                                                                                                                                                                                                                                                              [6]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Highly accurate genome seques MG1655 and W3110."; Mol. Syst. Biol. 0:0-0(2006)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Nucleotide sequence Escherichia coli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Smallshaw J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=K-12
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NUCLEOTIDE SEQUENCE
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NUCLEOTIDE SEQUENCE
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                                                                                                                NUCLEOTIDE
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 460-kb
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PubMed=8387990;
H., Feulner G.,
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RESULT 25

Q8X841_ECO57

ID Q8X841_ECO57 PRELIMINARY; PRT; 474 AA.

AC Q8X841; Q7ABC1;

DT 01-MAR-2002, integrated into UniProtKE/TrEMBL.

DT 01-MAR-2002, sequence version 1.

DT 07-FEB=2006, entry version 26.

DE 6-phospho-beta-glucosidase; cryptic (6-phospho-bet Name=ascB; OrderedLocusNames=EC83572, z4024;

OS Escherichia coli 0157:H7.

OC Bacteria; Proteobacteria; Gammaproteobacteria; Ent Enterobacteriaceae; Escherichia.

OC Riterobacteriaceae; Escherichia.

OX NCBI_TaxID=83334;

RN [1]

RP NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

RC STRAIN=0157:H7 / EDL933 / ATCC 700927 / EHEC;

RX MEDLINE=21074935; PubMed=11206551; DOI=10.1038/350

RA Perna N.T., Plunkett G. III, Burland V., Mau B., G.

RA Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kir

RA Posfai G., Hackett J., Klink S., Boutin A., Shaoy

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STRAIN=K-12;
STRAIN=S0094132; PubMed=1459951;
MEDLINE=93094132; PubMed=1459951;
Gervais F.G., Drapeau G.R.;
Gervais F.G., Cloning, and characterization of rcsF, a new
Gervais F.G., Cloning, and characterization of rcsF, a new
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STRAIN-0157:H7 / EDL933 / ATCC 700927 / HEIG.
MEDLINE-21074935; PubMed-11205551; DOI=10.1038/35054089;
Perna N.T., Plunkett G. III, Burland V., Mau B., Glasner J.D.,
Posfai G., Hackett J., Klink S., Gregor J., Kirkpatrick H.A.,
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Grotbeck E.J., Davis N.W., Link A., Dimalanta E.T., Potamousis K.,
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the mukB null mutant of Escherichia coli.";
J. Bacteriol. 174:7517-7526(1992).
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21-FEB-2006, entry version 4.
Permease for cytosine/purines, uracil,
Permease for enames=Pfl_3440;
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STRAIN-0157:H7 / Sakai / RIMD 0509952 / EHEG.

MEDLINE-21156231; PubMed=11258796; DOI=10.1093/dnares/8.1.11;

Hayashi T., Makino K., Ohnishi M., Kurokawa K., Ishii K., Yokoya
Han C.-G., Ohtsubo E., Nakayama K., Murata T., Tanaka M., Tobe 1

Iida T., Takami H., Honda T., Sasakawa C., Ogasawara N., Yasunag

Kuhara S., Shiba T., Hattori M., Shinagawa H.;

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0157:H7 and genomic comparison with a laboratory strain K-12.";
   NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

US DOE Joint Genome Institute;

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Larimer F., Land M., Kyripides N., Anderson I., Richardson
"Complete sequence of Pseudomonas fluorescens PfO-1.";

Submitted (AUG-2005) to the EMBL/GenBank/DDBJ databases.
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SEQUENCE
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BioCyc; ECOLB3334-1:ECS3572-MONOMER; -.
GO; GO:0004553; F:hydrolase activity, hydro
GO; GO:0004553; F:carbohydrate metabolism;
GO; GO:0005975; P:carbohydrate mydro_1.
InterPro; IPR001360; Glyco_hydro_1; 1.
PANTHER; PTHR10353; Glyco_hydro_1; 1.
Pfam; PF00232; Glyco_hydro_1; 1.
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                                                                          Copyrighted Distributed
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PROSITE; PS00572; GLYCOSYL_HYDROL_F1_1;
PROSITE; PS00653; GLYCOSYL_HYDROL_F1_2;
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EMBL; BA000007; BAB36995.1; -; Genomic_DNA.
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L; CP000094; ABA75178.1; -; Genomic_DNA. GO:0016020; C:membrane; IEA. GO:0015205; F:nucleobase transporter activity; GO:0015931; P:nucleobase, nucleoside, nucleotic
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10-JAN-2006,
Pseudomonas putida (встат
Bacteria; Proteobacteria;
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07-FEB-2006, entry version 10.
Transporter, NCS1 nucleoside transporter family.
OrderedLocusNames=PP4033; ORENames=PP4035;
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Larimer F., Land M.;
"Annotation of the draft genome assembly of Pseudomonas |
Submitted (NOV-2005) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
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Pfam; PF02133; Transp_cyt_pur; 1.
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Best Local
                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

STRAIN=ATCC 15692 / PAO1;

STRAIN=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINE=20437337; PubMed=10.1038/35023079;

MEDLINE=20437337; PubMed=10.984043; DOI=10.1038/35023079;

MEDLINE=20437337; PubMed=10984043; DOI=10.1038/35023079;

MEDLINE=20437337; PubMed=10.0038/35023079;

MEDLINE=2043737; PubMed=10.0038/35023079;

MEDL
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Nelson K.E., Weinel C., Paulsen I.T., Dodson R.J., Hilbert H.,
Martine dos Santos V.A.P., Fouts D.E., Gill S.R., Pop M., Holmes
Brinkac L.M., Beanan M.J., DeBoy R.T., Daugherry S.C., Kolonay J
Madupu R., Nelson W.C., White O., Peterson J.D., Khouri H.M.,
Hance I., Chris Lee P., Holtzapple E.K., Scanlan D., Tran K.,
Moazzez A., Utterback T.R., Rizzo M., Lee K., Kosack D., Moestl
Wedler H., Lauber J., Stjepandic D., Hoheisel J., Straetz M., He
Kiewitz C., Eisen J.A., Timmis K.N., Duesterhoeft A., Tuemmler B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q91674;
01-MAR-2001, integrated into UniProtKB/TrEMBL.
01-MAR-2001, sequence version 1.
07-FBB-2006, entry version 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL;
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                                 EMBL;
                                                                                                                            Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                   "Complete genome sequence of Pseudomonas aeruginosa opportunistic pathogen.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteria; Proteobacteria; Gammaproteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OrderedLocusNames=PA0443;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Probable transporter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fraser C.M.;
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NCBI_TaxID=160488;
                                                                                                                                                                                                                                                                                 Nature 406:959-964(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=287;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pseudomonadaceae;
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GO:0015205; F:nucleobase transporter activity; IE
GO:0015931; P:nucleobase, nucleoside, nucleotide
; AE004481; AAG03832.1;
B83591; B83591.
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Pred. No. 1.3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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blonay J.
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Lim R.M.,
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RESULT 31
QBPQLB XANAC
ID GBPQLB X
AC QBPQLB,
DT 01-OCT-2
DT 01-CCT-2
DT 07-FEB-2
DT 07-FEB-2
DE Gamma-91
GN Name=99t
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Q3BYW8;
22-NOV-2005,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

PubMed=16237009; DOI=10.1128/JB.187.21.7254-66.2005;

Phieme F., Koebnik R., Bekel T., Berger C., Boch J., Buettner Caldana C., Gaigalat L., Goesmann A., Kay S., Kirchner O., La Linke B., McHardy A.C., Meyer F., Mittenhuber G., Nies D.H., Niesbach-Kloesgen U., Patschkowski T., Rueckert C., Rupp O., Schneiker S., Schuster S.C., Vorhoelter F.J., Weber E., Puehl Bonas U., Bartels D., Kaiser O.;

"Insights into genome plasticity and pathogenicity of the pla pathogenic Bacterium Xanthomonas campestris pv. vesicatoria r by the complete genome sequence.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BioCyc; PAER287:PA0443-MONOMER; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0015205; F:nucleobase transporter activity; IEA.
GO; GO:0015331; P:nucleobase, nucleoside, nucleotide and nucl.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AM039952; CAJ21945.1; -; Genomic_DNA.
GO; GO:0008415; F:acyltransferase activity; IEA.
GO; GO:0003840; F:gamma-glutamyltransferase activity;
GO; GO:0016740; F:transferase activity; IEA.
Acyltransferase; Complete proteome; Transferase.
SEQUENCE 526 AA; 55499 MW; B40B3B74381BAA43 CRC64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Xanthomonas campestris pv. vesicatoria (strain 85-10).
Bacteria; Proteobacteria; Gammaproteobacteria; Xanthomonadales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-NOV-2005,
07-FEB-2006,
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Pfam; PF02133; Transp_cyt_pur; 1.
                                                  01-OCT-2002, integrated into t
01-OCT-2002, sequence version
07-FEB-2006, entry version 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  J. Bacteriol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=ggt1; OrderedLocusNames=XCV0314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gamma-glutamyltranspeptidase (EC 2.3.2.2).
                          Gamma-glutamyltranspeptidase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             complete genome sequence.";
teriol. 187:7254-7266(2005).
                                                                                                                                                     XANAC
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Pred. No. 1.4
0; Mismatches
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Pred. No.
                                                                                                                                                     PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B40B3B743818AA43 CRC64;
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RA Quaggio R.B., Monteiro-Vitorello C.B., Van Sluys M.A.,
RA Quaggio R.B., Monteiro-Vitorello C.B., Van Sluys M.A.,
RA Almeida N.F. Jr., Alves L.M.C., do Amaral A.M., Bertolini M.C.,
RA Camargo L.B.A., Camarotte G., Cannavan F., Cardozo J., Chambergo F.,
RA Ciapina L.P., Cicarelli R.M.B., Coutinho L.L., Cursino-Santos J.R.,
RA El-Dorry H., Faria J.B., Ferreira A.J.S., Ferreira R.C.C.,
RA Ferro M.I.T., Formighieri E.F., Franco M.C., Greeggio C.C., Gruber A.,
RA Katsuyama A.M., Kishi L.T., Leite R.P., Lemos E.G.M., Lemos M.V.F.,
RA Locali E.C., Machado M.A., Madeira A.M.B.N., Martinez-Rossi N.M.,
RA Martins E.C., Meidanis J., Menck C.F.M., Miyaki C.Y., Moon D.H.,
RA Moreira L.M., Novo M.T.M., Okura V.K., Oliveira M.C., Oliveira V.R.,
RA Pereira H.A., Rossi A., Sena J.A.D., Silva C., de Souza R.F.,
RA Spinola L.A.F., Takita M.A., Tamura R.E., Teixeira B.C., Tezza R.I.D.,
RA Setubal J.C., Kitajima J.P.;
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Matches 6
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29 NEUCR
Q7SA29 NEUCR
Q7SA29;
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15-DEC-2003, sequence version 1.
07-FEB-2006, entry version 11.
Hypothetical protein.
ORFNames=NCU07334.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GO; GO:0003840; P:gamma-glutamyltransferase InterPro; IPR000101; GGT peptidase. PANTHER; PTHR11686; GGT_peptidase; 1. Pfam; PF01019; G glu transpept; 1. PRINTS; PR01210; GGTRANSPTASE.
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STRAIN=74-OR23-1A / FGSC 987,
MEDLINE=22598136; PubMed=12712197; DOI=10.1038/nature01554;
MEDLINE=22598136; PubMed=12712197; DOI=10.1038/nature01554;
Galagan J.E., Calvo S.E., Borkovich K.A., Selker E.U., Read N.D.,
Jaffe D., FitzHugh W., Ma L.-J., Smirnov S., Purcell S., Rehman B
Elkins T., Engels R., Wang S., Nelsen C.B., Butler J., Endrizzi
Qui D., Ianakiev P., Bell-Pedersen D., Nelson M.A.,
Werner-Washburne M., Selitrennikoff C.P., Kinsey J.A., Braun E.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Xanthomonadaceae; Xanthomonas.
                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Fungi; Ascomycota; P
Sordariomycetidae; Sordariales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neurospora crassa.
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Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nature 417:459-463(2002).
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                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE [LARGE
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                                                                                                                                                                                                                                                                                                                                                                                           Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., G Hammon N., Israni S., Pitluck S., Chain P., Malfatti S., Sh Vergez L., Schmutz J., Larimer F., Land M., Hauser L., Pell Kyrpides N., Anderson I., Oda Y., Harwood C.S., Richardson "Complete sequence of Rhodopseudomonas palustris HaA2."; Submitted (JAN-2006) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Extracellular solute-binding ORFNames=RPB_1770; Rhodopseudomonas palustris Ha
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07-MAR-2006, sequence version 1.
07-MAR-2006, entry version 1
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GO; GO:0015205; F:nucleobase transport
GO; GO:0015931; P:nucleobase, nucleosi
InterPro; IPR001248; Cyt_pur_permease.
InterPro; IPR012681; NCS1.
Pfam; PF02133; Transp_cyt_pur; 1.
TIGRFAMs; TIGR00800; ncs1; 1.
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NCBI_TaxID=316058;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "The genome sequence of the filamentous fungus Neurospora crassa."; Nature 422:859-868(2003).
                                                                                                                                                                                                                        EMBL;
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SEQUENCE 50
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CAUTION: The sequence shown here is derived from an
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D:0016020; C:membrane; IEA.
                                                                                                                                                                                                                        CP000250;
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560 AA;
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F:nucleobase transporter activity; IEA.
                                                                                                                                                                                                                        ABD06478.1; -;
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Pred. No. 1.5e+02;
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Pelletier D.A.,
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Q4SCJ9;
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"Annotation of the draft genome of Phoses.";
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Q374J7;
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SIGNAL
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Copeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
"Sequencing of the draft genome and assembly
palustris BisA53.";
                                                                                                                                                                               19-JUL-2005, integrated into UniProtKB/TrEMBL.
19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 6.
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                            Tetraodon nigroviridis (Green Eukaryota; Metazoa; Chordata;
                                                                                 (Fragment).
ORFNames=GSTENG00020461001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                                                         SCAF14653,
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n puffer).
; Craniata;
Teleostei;
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32F26A217EC705E4 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.8e+02;
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Euteleostei;
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                               Euteleostomi;
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RESULT 36
Q4P6R1 USTWA
ID Q4P6R1 U
AC Q4P6R1;
DT 19-UUL-2
DT 19-UUL-2
DT 19-UUL-2
DT 07-FEB-2
DE Hypothet
GN ORFNames
OS Ustilago
OC Eukaryot
OC Ustilago
OX NCBI_Tax
RN [1]
RP NUCLEOTI
RC STRAIN=5
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RA Jaillon O., Aury J.-M., Brunet F., Petit J.-L., Stange-Thomann N.,
RA Jaillon O., Aury J.-M., Brunet F., Petit J.-L., Stange-Thomann N.,
RA Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
RA Micaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Anthouard V., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
RA Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,
RA Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,
RA Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
RA Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA Lindblad-Toh K., Schachter V., Quetier F., Saurin W., Scarpelli C.,
Mincher P., Lander E.S., Weissenbach J., Roest Crollius H.;
"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
The early vertebrate proto-karyotype.";
U. Nature 431:946-957 (2004).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                    19-JUL-2005,
19-JUL-2005,
07-FEB-2006,
                                                                                                                                                                                                                        Hypothetical protein. ORFNames=UM04202.1;
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Q4P6R1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genoscope; Whitehead Institute Centre for Genome Research; Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
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   NUCLEOTIDE
STRAIN=521;
                                                                                                                              Eukaryota; Fungi; Basidiomycota; Us
Ustilaginomycetidae; Ustilaginales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS01013; OSBP; 1.
PROSITE; PS50003; PH_DOMAIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF01237; Oxysterol BP; PRINTS; PR00360; C2DOMAIN.
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Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                          Ustilago maydis 521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR000008; C2.
InterPro; IPR000648; Oxysterol_bd.
InterPro; IPR001849; PH.
                                                                                             NCBI_TaxID=237631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    L; CAAE01014653; CAG01633.1; -; Genomic D
GO:0003676; F:nucleic acid binding; IEA.
GO:0006869; P:lipid transport; IEA.
GO:0008202; P:steroid metabolism; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 preliminary data.
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ER 809 809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FSRYAR 747
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                                    SEQUENCE
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                                                                                                                                                                                                                                                                                       entry version 4.
                                                                                                                                                                                                                                                                                                                                                   integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
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                                    [LARGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 31; DB 2; 100.0%; Pred. No. 2.2e+02; tive 0; Mismatches 0;
                                    SCALE
                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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                                                                                                                       Ustilaginomycetes;
es; Ustilaginaceae; Ustilago.
                                    GENOMIC
                                                                                                                                                                                                                                                                                                                                                                                                                  1427
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                                                                                                                                   Matches
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Best Local
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InterPro; IPR006615; Pept C19 N 1.
InterPro; IPR001394; Peptidase_C19.
Pfam; PP00443; UCH; 1.
SMART; SM00695; DUSP; 1.
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                                                                                                                                                                                                                                                                                                                       PROSITE; PS00147; ARGINASE 1; UNKNOWN 1
PROSITE; PS00972; UCH 2 1; 1.
PROSITE; PS00973; UCH 2 2; 1.
PROSITE; PS50235; UCH 2 3; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "The genome sequence of Ustilago maydis.";
Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                                                                                                                                                                                                                                                               Hypothetical SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GO; GO:0004197; F:cysteine-type endopeptidase activity; GO; GO:0004221; F:ubiquitin thiolesterase activity; IEA GO; GO:0006512; P:ubiquitin cycle; IEA GO; GO:0006512; P:ubiquitin cycle; IEA GO; GO:0006511; P:ubiquitin-dependent protein catabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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896 FSRYAR 901
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                                                                                                                               Similarity 6; Conserv
                                                                                                                                                                                                                                                                  ni protein.
1427 AA;
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Pred. No. 4e+02;
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DT 01-QC
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GO; GO:0003887; F:DNA-directed DNA polymerase activity; 1
GO; GO:0006260; F:DNA replication; IEA.
InterPro; IPR001998; DNA pol.
InterPro; IPR002298; DNA polI.
Pfam; PF00476; DNA polI A; 1.
PRINTS; PR00868; DNAPOLI.
PROSITE; PS00447; DNA POLYMERASE A; UNKNOWN 1.
SEQUENCE 1649 AA; 182029 MW; 2FC69472CCZA7A1E CRC64;
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01-OCT-2002, sequence version 1.
07-FEB-2006, entry version 15.
DNA polymerase I-like protein C.
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Bukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae; Trypanosoma.
                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                           Klingbeil M.M., Motyka S.A., Englund P.T.;
"Multiple mitochondrial DNA polymerases in Trypanosoma brucei.";
Mol. Cell 10:175-186(2002).
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                                      L; AF445378; AAM81964.1; -; Genomic_DNA. GO:0003677; F:DNA binding; IEA. GO:0003887; F:DNA-directed DNA polymerase GO:0006260; P:DNA replication; IEA.
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10-MAY-2005, sequence version 1.
07-FEB-2006, entry version 2.
Hypothetical protein PHG310RF237c.
Name=PHG310RF237c; ORFNames=PHG31p234;
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PRINTS; PRO0868; DNA POLYMERASE A; UNKNOWN 1.
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PROSITE; PS00447; DAABC43C737A3BB0 CRC64;
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"The gyrB gene lies opposite from circular chromosome of Mycoplasma
                                                                                                                                                                                                                                                                      Distributed under the Creative Commons Attribution-NoDerivs
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Sano K., Miyata M.;
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 Viruses; dsDNA viruses,
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"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";

Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
Actinopterygii; Neopterygii; T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                              Copyrighted
Distributed
                                                                                                                                                                                                                        Director MGC Project; Submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                          TISSUE=Larvae;
Director MGC E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Larvae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
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SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=wrb;
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                                 Ensembl;
                                                               EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=7955;
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BRARE
                                                               BC085456; AAH85456.1; -; mRNA.
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   ZDB-GENE-030131-7696;
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5; Conserv
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                              ENSDARG00000024641; Danio rerio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            al protein.
102 AA; 1
                                                                                                                        by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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83.3%;
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Pred. No. 1.3e
1; Mismatches
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Teleostei;
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Craniata; Vertebrata; Euteleostomi;
Teleostei; Ostariophysi; Cypriniform
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hage 31 genome.";
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RESULT 43
Q3JOBB BURP1 PRELIMINARY; PRT; 133 AA.
ID Q3JOBB_BURP1 PRELIMINARY; PRT; 133 AA.
AC Q3JOBS;
DT 08-NOV-2005, integrated into UniProtKB/TrEMBL.
DT 08-NOV-2005, sequence version 1.
DT 21-FEB-2006, entry version 4.
DT 21-FEB-2006, entry version 4.
DE Hypothetical protein.
GN OrderedLocusNames=BURPS1710b 2850;
OS Burkholderia pseudomallei (scrain 1710b).
OC Bacteria; Proteobacteria; Betaproteobacteria; Bur OC Burkholderiaceae; Burkholderia; pseudomallei grou OX NCBI TaxID=320372;
RN UNCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
RA Woods D.E., Nierman W.C.;
RL Submitted (SEP-2005) to the EMBL/GenBank/DDBJ dat
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                                                                                                                                                                                                                                                                                                                                                                                                    Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Caldwell R.B., Kierzek A.M., Arakawa H., Bezzubov Y., Zaim J., Fiedler P., Kutter S., Blagodatski A., Kostovska D., Koter M., Plachy J., Carrinci P., Hayashizaki Y., Buerstedde J.M., "Full-length cDNAs from chicken bursal lymphocytes to facilita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-NOV-2004, sequence version 07-FEB-2006, entry version 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q5ZJM2_CHICK
Q5ZJM2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR007514; CHD5
Pfam; PF04420; CHD5; 1.
                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein. SEQUENCE 126 AA; 1
                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR007514; CH Pfam; PF04420; CHD5; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AJ720412; CAG32071.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genefunction analysis."
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORFNames=RCJMB04
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CHICK
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FARYAR 10
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     EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                    Score 28;
Pred. No.
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Pred.
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Attribution-NoDerivs License
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                                                                                                                                                                                                   XX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
XS Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
XA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
XA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
XA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
XA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
XA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
XA Roman S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
XA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
XA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
XA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
XA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
XA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
XA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
XA Holder J., Schuntz J., Myers R.M.,
XA Holder J., Schein J.E., Jones S.J.M., Marra M.A.,

"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.",

171 Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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Hypothetical protein.
Xenopus laevis (African clawed frog).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q52KS6_XENLA
Q52KS6;
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                                                          Submitted (APR-2005) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=22341132; PubMed=12454917; DOI=10.1002/dvdy.10174; Klein S.L., Strausberg R.L., Wagner L., Pontius J., Clifton S.W.,
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SEQUENCE 133 AA; 14144 MW; C069DFC776D45162 CRC64;
                                                                                                                                                    NUCLEOTIDE SEQUENCE
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                                                                                                                        rissue=Eye;
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RESULT 46
Q5AAF5 CANAL
ID Q5AAF5 CANAL
AC Q5AAF5;
AC Q5AAF5;
DT 26-APR-2
DT 26-APR-2
DT 97-FEB-2
DT Hypothet
GN ORFNames
OS Candida
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InterPro; ...
InterPro; ...
170 f
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                                                                                             Candida albicans SC5314.
Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.

PubMed=15256591; DOI=10.1073/pnas.0403929101;

PubMed=15256591; Doi=10.1073/pnas.0403929101;

Nissen R.M., Sun Z., Swindell E.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata, Actinopterygii; Neopterygii; Cyprinidae; Danio.
NCBI_TaxID=7955;
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STRAIN=SC5314;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Copyrighted by the UniProt Consortium, see http://www.uniprot.org/to
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Eukaryota; Metazoa; Chordata;
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                                                                    NCBI_TaxID=237561;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         development.";
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Pred. No. 2.2e+02;
                                                                                             Saccharomycotina; Saccharomycetes; Saccharomycetales; Candida.
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RA Park H.-S., Toyoda A., Ishii K., Totoki Y., Choi D.-K., Groner Y.,
RA Sceda E., Ohki M., Takagi T., Sakaki Y., Taudien S., Blechschmidt K.,
RA Polley A., Menzel U., Delabar J., Kumpf K., Lehmann R., Patterson D.,
RA Reichwald K., Rump A., Schillhabel M., Schudy A., Zimmermann W.,
RA Rosenthal A., Kudoh J., Shibuya K., Kawasaki K., Asakawa S.,
RA Rosenthal A., Sasaki T., Nagamine K., Mitsuyama S., Antonarakis S.E.,
RA Minoshima S., Shimizu N., Nordsiek G., Hornischer K., Brandt P.,
RA Scharfe M., Schoen O., Desario A., Reichelt J., Kauer G., Bloecker H.,
RA Romser J., Beck A., Klages S., Hennig S., Riesselmann L., Dagand E.,
RA Wehrmeyer S., Borzym K., Gardiner K., Nizetic D., Francis F.,
RA Lehrach H., Reinhardt R., Yaspo M.-L.;
RT "The DNA sequence of human chromosome 21.";
LNature 405:311-319(2000).
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"The diploid genome sequence of Candida albicans.";
Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334(2004).
-I- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WRB HUMAN STANDARD; PRT; 174 AA. 000258; 060740; 01-NOV-1997, integrated into UniProtKB/Swiss-Prot. 01-DEC-2000, sequence version 2. 07-FEB-2006, entry version 34. Tryptophan-rich protein (Congenital heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Egeo A., Mazzocco M., Sotgia F., Arrigo P., Oliva R., Bergono Nizetic D., Rasore-Quartino A., Scartezzini P.; "Identification and characterization of a new human cDNA from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PubMed=15123810; DOI=10.1073/pnas.0401648101;
Jones T., Federspiel N.A., Chibana H., Dungan
Magee B.B., Newport G., Thorstenson Y.R., Agal
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
MEDLINE=20289799, PubMed=10830953; DOI=10.1038/35012518;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 21q22.3 encoding a basic nuclear protein.";
Hum. Genet. 102:289-293(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=WRB; Synonyms=CHD5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AACQ01000039; EAK99775.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=98204400; PubMed=9544840; DOI=10.1007/s004390050693;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            preliminary data.
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5; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ai protein.
171 AA; 1
                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17791 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    [MRNA],
                                                                                             [LARGE SCALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.3%;
83.3%;
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Pred. No. 2.2e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                032E48F64D7C8B2E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION
                                                                                             MRNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        heart disease 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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RESULT
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RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
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RA Rosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Hichards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOUSE
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SEQUENCE
STRAIN=C57BL/6J; TISSUE=Corpus striatum;
PubMed=16141072; DOI=10.1126/science.1112014;
Carninci P., Kasukawa T., Katayama S., Gough J., Frith M.C., Maeda I Oyama R., Ravasi T., Lenhard B., Wells C., Kodzius R., Shimokawa K. Bajic V.B., Brenner S.E., Batalov S., Forrest A.R., Zavolan M., Davis M.J., Wilming L.G., Aidinis V., Allen J.B.,
Ambesi-Impiombato A., Apweller R., Aturaliya R.N., Bailey T.L.,
Bansal M., Baxter L., Beisel K.W., Bersano T., Bono H., Chalk A.M.,
Chiu K.P., Choudhary V., Christoffels A., Clutterbuck D.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WRB_MOUSE STANDARD; PRT; 174 AA.

QBKOD7; Q9D1W0;

Q0-MAY-2005, integrated into UniProtKB/Swiss-Prot.

Q1-OCT-2002, sequence version 1.

Q7-FEB-2006, entry version 18.

Tryptophan-rich protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; Y12478; CAA73081.1; -; mRNA.
EMBL; AF064861; -; NOT ANNOTATED_CDS; Genomic_EMBL; AL163279; CAB90454.1; -; GENOMIC_DNA.
EMBL; BC012415; AAH12415.1; -; mRNA.
ENSEMD1; ENGG00000182093; Homo sapiens.
H-InvDB; HIX0016117; -.
                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA]
                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                 Muroidea;
                                                                                                                                                                                                                                                                                                                                                                                              Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nuclear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF04420; CHD5; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; 602915; gene.
GO:0005634; C:nucleus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             c. Natl. Acad. Sci. U.S.A. 99:10
SUBCELLULAR LOCATION: Nucleus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           69
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5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein.
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FARYAR
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Muridae; N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31
174 AA;
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                                                                                                                                                                                                                                                                                                                                                                     Murinae;
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19780
                                                                                                                                                                                                                                                                                                                                                                                              Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurogna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHD5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A. 99:16899-16903(2002).
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                                                                                                                                                                                                                                                                                                                                                                     Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 28;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tryptophan-rich protein.

/FTId=PRO_0000065979.

F -> I (in Ref. 1).

; A01F11CC564EAB6E CRC64
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RA Gustincich S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa N., Ra Hill D., Huminiecki L., Tacono M., Ikeo K., Iwama A., Ishikawa T., Ra Hill D., Huminiecki L., Tacono M., Kelso J., Kitamura H., Ra Jakt M., Kanapin A., Katoh M., Kawasawa Y., Kelso J., Kitamura H., Ra Kitano H., Kollias G., Krishnan S.P., Kruger A., Kummerfeld S.K., Ra Kurochkin I.V., Lareau L.F., Lazarevic D., Lipovich L., Liu J., Ra Kitano H., McHiliam S., Madan Babu M., Madera M., Marchionni L., Ra Kurochkin I.V., Lareau L.F., Lazarevic D., Lipovich L., Liu J., Ra Mctuda H., Mateuzawa S., Miki H., Mignone F., Miyake S., Morris K., Ra Milson R., Nishiguchi S., Nishikawa S., Nori F., Ohara O., Olardo V., Paug K.C., Pavan W.J., Pavesi G., Pesole G., Petrovsky N., Piazza S., Reed J., Reid J.F., Ring B.Z., Ringwald M., Rost B., Ruan Y., Salzberg S.L., Sandelin A., Schneider C., Schipachi K., Semple C.A., Seno S., Sessa L., Sheng Y., Shibata Y., Shimada H., Shimada K., Silva D., Sinclair B., Sperling S., Stupka E., Sugiura K., Sultana R., Takenaka Y., Taki K., Anamoja K., Tan S.L., Tang S., Taylor M.S., Tegner J., Teichmann S.A., Veda H.R., van Nimwegen E., Verardo R., Wei C.L., Yagi K., Yananishi H., Zabarovsky E., Zhu S., Zimmer A., Hide W., Bult C., Ra Kawashima T., Kojima M., Kondo S., Konno H., Makano K., Ninomiya N., Ra Tida J., Tamura K., Itoh M., Kato T., Kawaji H., Kawagashira N., Ra Ra Nishio T., Okada M., Plessy C., Shibata K., Shiraki T., Suzuki S., Tagami M., Waki K., Watahiki A., Okamura-Oho Y., Suzuki H., Kawai J., Ra Hayashizaki Y..
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Vallalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Vallalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Vallalon D.K., Wadan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Miting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Rodriguez A.C., Grinwood J., Schmutz J., Myers R.M.,
A Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
"Generation and initial analysis of more than 15,000 full-length human
"Tennewise C.N.M. Generation and initial analysis of more than 15,000 full-length human
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di Bernardo
Fletcher C.F
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[2]
                                        GO;
                                                                             EMBL; AK021091; BAB32297.1; -; mRNA. EMBL; BC031769; AAH31769.1; -; mRNA. Ensembl; ENSMUSG00000003147; Mus musmg1; mGI:2136882; Wrb.
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                                                                                                                                                                                                                                                                                                   Copyrighted
Distributed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proc.
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GO:0016021; C:integral to membrane; TAS erPro; IPR007514; CHD5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mouse cDNA sequences.";
c. Natl. Acad. Sci. U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Nucleus ALTERNATIVE PRODUCTS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Event=Alternative
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                                                                                                                                                                                                                                                                                                                                                                                                                                  IsoId=Q8K0D7-2; Sequence=VSP_013643, VSP_013644; \
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IsoId=Q8K0D7-1; Sequence=Displayed;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rdo D., Down T., Engstrom P., Fagiolini M., Faulkner G., C.F., Fukushima T., Furuno M., Futaki S., Gariboldi M., Henming P., Gingeras T.R., Gojobori T., Green R.E., Ch. S., Harbers M., Hayashi Y., Hensch T.K., Hirokawa N.,
                                                                                                                                                                                                                                                              by the UniProt Consortium, see http://www.uniprot.org/terms
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D., Down T., Engstrom P., Fagiolin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      splicing; Named
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WRB_RAT
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05P6S5;

10-MAY-2005, integrated into U

05-JUL-2004, sequence version

07-FEB-2006, entry version 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QSRGKY;
10-MAY-2005, integrated into UniProtKB/Swiss-Prot.
21-DEC-2004, sequence version 1.
07-FEB-2006, entry version 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WRB
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Alternative splicing; Nuclear CHAIN 1 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pongo pygmaeus (Orangutan).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                             Rattus norvegicus (Rat).
                                                                                    Tryptophan-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR007514; CH
Pfam; PF04420; CHD5; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; CR860481; CAH92603.1; -; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The German cDNA consortium; Submitted (NOV-2004) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tryptophan-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurogna
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Pred. No.
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Pred. No.
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Missing (in isoform 2)
/rmrd=VSP 013644.
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/FTId=PRO_000065981.
; OF1F0F3DB1D38202 CRC64;
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/FTId=PRO_0000065980.
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2.3e+02;
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Ensembl; ENSRNOG0000001629; Rattus norvegicus.
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InterPro; IPR007514; CHD5.
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69 FARYAR 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 174 Tryptophan-rich protein.
/FTId=PRO_000065982.
174 AA; 19899 MW; 7941B91831F2DC6B CRC64;
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sequence 11967, A Sequence 20570, A Sequence 464, App Sequence 15610, A Sequence 994, App Sequence 218, App Sequence 218, App Sequence 10670, A Sequence 7196, Ap Sequence 3, Appli Sequence 3, Appli Sequence 7362, Ap		Sequence 17,, App Sequence 154, App Sequence 45424, A Sequence 2553, A Sequence 50, Appl	Sequence 46444, A Sequence 3129, Ap Sequence 61958, A Sequence 14484, A Sequence 17, Appl	Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 43460, A Sequence 18, Appl	Sequence 48109, A Sequence 1, Appli Sequence 3921, Ap Sequence 3921, Ap	Sequence 15.28, A Sequence 26, Appl Sequence 28, Appl Sequence 14040, A Sequence 32892. A	Sequence 5491, Ap Sequence 17800, A Sequence 48, Appl Sequence 48, Appl	Sequence 102, App Sequence 467, App Sequence 10579, A Sequence 10579, A	Sequence 38, Appl Sequence 38, Appl Sequence 38, Appl	Sequence 13913, Ap Sequence 5171, Ap Sequence 13903, A	Sequence 5490, Ap Sequence 19, Appl Sequence 19, Appl	Sequence 35068, A Sequence 50285, A Patent No. 5320958	857, Ā	3, App	8, App	i mi	60533, 1300, A
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US-09-252-991A-28081
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US-09-526-3098-3
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; Sequence 4, Application US/08928862
; Patent No. 6309877
; GENERAL INFORMATION:
; APPLICANT: Chau, Raymond M. W.
; TITLE OF INVENTION: Isolation and Use of Mot
; FILE REFERENCE: 12592-2
; CURRENT APPLICATION NUMBER: US/08/928,862
; CURRENT FILING DATE: 1997-09-12
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 4
; LENGTH: 33
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-928-862-4
 US-09-592-018-4
; Sequence 4, Application US/09592018
; Patent No. 6759389
; GENERAL INFORMATION:
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US-08-587-389-1
US-08-587-389-2
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US-09-653-274-10
US-09-302-769-44
US-09-712-363-180
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RESULT 4
US-09-198-452A-977
; Sequence 977, Application US/09198452A
; Patent No. 6559294
; GENERAL INFORMATION:
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; ORGANISM: Homo Sapiens
US-09-989-481-4
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US-09-989-481-4
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US-09-592-018-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/633,447
PRIOR FILLING DATE: 2000-08-07
PRIOR PELLY DATE: 1097-09-12
PRIOR PILLING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILLING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 66/026792
PRIOR APPLICATION NUMBER: US 66/026792
PRIOR FILLING DATE: 1996-09-27
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-4
APPLICANT: Griffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragment:
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, prevention of INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/989,481
CURRENT FILING DATE: 2001-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/026792
PRIOR FILING DATE: 1996-09-27
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/592,018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 12592-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 33
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APPLICANT: The Regents of the University of California TITLE OF INVENTION: Chlamydia pneumoniae Genome Sequence FILE REFERENCE: 018941-000411US
CURRENT APPLICATION NUMBER: US/09/438,185A
CURRENT FILING DATE: 2002-03-13
PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR FILING DATE: 1998-11-12
PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
SOFTWARE: FASTEEQ for Windows Version 3.0
SEQ ID NO 906
LENGTH: 359
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US-09-438-185A-906
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CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 977
LENGTH: 357
                                                                                                                                                                                                                                                                                                                                                                                                               Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Stephens, Richard
APPLICANT: Mitchell, Wayne
APPLICANT: Kalman, Sue
                                                                                                                                                                                                                   Sequence 60, Application US/08914375C Patent No. 6377893 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 906, Application US/09438185A Patent No. 6822071
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TYPE: PRT
ORGANISM: Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Chlamydia pneumoniae
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
              ZIP: 32605-4147
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                             Applications of Protein Structure Predictions
NUMBER OF SEQUENCES: 74
                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                            137 FSRYAR 142
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                                                                                                                                                                                                                                                                                                                                                                              1 FSRYAR 6
                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 6; Conserv
                                                                                    STATE:
                                                                                                     CITY: Gainesville
                                                                                                                 ADDRESSEE: Steven A. Benner STREET: 1501 NW 68th Terrace
COMPUTER: Apple MacIntosh
                                                                 COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Davis, Ronald
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100.0%; Pred. No. (
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; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-23341
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US-09-252-991A-23341
; Sequence 23341, Application US/09252991A
; Patent No. 6551795.
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GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION UNMEER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US (
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 23341
LENGTH: 509
                                                                                                                                   Sequence 11070, Application US/09949016
Patent No. 6812339
                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Best Local
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: amino acid
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/914,375C
FILING DATE: 19-Aug-1997
CLASSIFICATION: 702/20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 352 392 7773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Microsoft Word CURRENT APPLICATION DATA:
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ilarity 100.0%;
Conservative (
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BER: US 60/074,788
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Pred. No. 82;
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US-09-252-991A-32051
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US-09-252-991A-18159
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SEQ ID NO 11070
LENGTH: 188
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LENGTH: 1778
TYPE: PRT
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GENERAL INFORMATION:
            NUMBER OF SEQ ID NOS:
SEQ ID NO 32051
                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                          Sequence 32051, Application US/09252991A Patent No. 6551795
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Best Local :
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Best Local
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                                                    CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788 PRIOR FILING DATE: 1998-02-18 PRIOR APPLICATION NUMBER: US 60/094,190 PRIOR FILING DATE: 1998-07-27
                                                                                                                                                                          TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 107196.136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
PRIOR FILING DATE: 1998-07-27
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/231,498 PRIOR FILING DATE: 2000-09-08
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
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ENGTH: 198
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1131 FSKYAR 1136
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5; Conserv
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5; Conserve
                                                                                                                                                                                                                                       Marc J.
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NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
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Pred. No. 1.4e+03;
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                   FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: 0.09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-9
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 14027
                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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GENERAL INFORMATION:
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                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 6610836
                                                                                                                                                                                                                                                                                                          APPLICANT: GATY Breton et. al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 2709.2004001
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 14027,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
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                                                                                                                                              LENGTH: 387
TYPE: PRT
ORGANISM: Klebsiella pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Pseudomonas aeruginosa
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nes 5; Conserv
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111 FSRYAQ 116
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83.3%;
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Pred. No. 2.4e+02;
1; Mismatches 0
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Pred. No. 4.5e+02;
                                                           DB 2; Le..
. 4.8e+02;
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RESULT 13

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; OTHER INFORMATION: arbb erwch; SEQUENCE DESCRIPTION: SEQ ID NO: 61: US-08-914-375C-61
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; Sequence 638, Application US/09198452A
; Patent No. 6559294
; GENERAL INFORMATION:
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                                                                                                                   ; ORGANISM: Chlamydia
US-09-198-452A-638
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OPERATING SYSTEM: MacIntosh 7.0
SOFTWARE: Microsoft Word
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/914,375C
PILING DATE: 19-Aug-1997
CLASSIFICATION: 702/20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 352 392 7773
TELEPHONE: 352 331 0462
INFORMATION FOR SEQ ID NO: 61:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                  SEQ ID NO 638
LENGTH: 460
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GENERAL INFORMATION:
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                                                  Matches
                                                               Query Match
Best Local
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TITLE OF INVENTION: Chamydia pneumoniae genomic sequence
TITLE OF INVENTION: thereof and uses thereof, in particul
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
                                                                                                                                                      TYPE: PRT
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: amino acid
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Applications of Protein Structure Predictions NUMBER OF SEQUENCES: 74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      150 FERYAR 155
1 FSRYAR 6
                                               Similarity
5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Steven A. Benner STREET: 1501 NW 68th Terrace
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83.3%;
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Pred. No. 5
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Pred. No. 5.
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5.7e+02;
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                                                                   .7e+02;
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particular for the diagnosis,
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; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 8224
; LENGTH: 469
; TYPE: PRT
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US-10-026-140-2
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US-09-489-039A-8224
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                                                                          GENERAL INFORMATION:
APPLICANT: Dunn-Coleman, Nigel
APPLICANT: Goedegebuur, Frits
APPLICANT: Ward, Michael
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                                                                                                                                                      Sequence 2, Application US/10026140 Patent No. 7005289
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APPLICANT: Gary Br
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CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gary Breton et. al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: 2709.2004001
APPLICANT: Yao, Jian
TITLE OF INVENTION: BGL5 Beta-Glucosidase and Nucleic Acids
TITLE OF INVENTION: Encoding the Same
FILE REFERENCE: GC697
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ORGANISM: Klebsiella pneumoniae
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                                                                        Ward,
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83.3%;
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Pred. No. 5.7e+02;
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Pred. No.
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5.9e+02;
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; ORGANISM: Trichoderma reesei; FEATURE: ; NAME/KEY: VARIANT; LOCATION: (1)...(484); OTHER INFORMATION: Xaa = Any Amino US-10-026-140-2
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US-09-134-001C-4633
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LENGTH: 484
TYPE: PRT
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SEQ ID NO 4633
LENGTH: 538
TYPE: PRT
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APPLICANT: The Regents of the University of California TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence FILE REFERENCE: 018941-000411US
CURRENT APPLICATION NUMBER: US 09/438,185A
CURRENT FILING DATE: 2002-03-13
PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR FILING DATE: 1998-11-12
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
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CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 3
                                                                                                                                                                APPLICANT: Stephens, Richard
APPLICANT: Mitchell, Wayne
APPLICANT: Kalman, Sue
APPLICANT: Davis, Ronald
APPLICANT: The Regents of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERWIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR PELING DATE: 1997-11-08
PRIOR PELING DATE: 1997-08-14
PRIOR FILING DATE: 1997-08-14
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83.3%;
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Pred. No.
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RESULT 20
US-09-602-787A-548
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SEQ ID NO 599
LEUGTH: 579
TYPE: PRT
ORGANISM: Chlamydia pneumoniae
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PRIOR FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/602,787A
CURRENT FILING DATE: 2000-06-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING PROTEINS
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OR FILING DATE: 1999-07-09
OR APPLICATION NUMBER: DE 19932124.8
OR FILING DATE: 1999-07-09
OR APPLICATION NUMBER: DE 19932125.6
OR FILING DATE: 1999-07-09
OR APPLICATION NUMBER: DE 19932128.0
OR APPLICATION NUMBER: DE 19932128.0
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APPLICATION NUMBER: DE 19931478.0
FILING DATE: 1999-07-08
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APPLICATION NUMBER: DE 19932227.9
FILING DATE: 1999-07-09
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                  APPLICATION NUMBER: DE 19932927.3 FILING DATE: 1999-07-14
                                                                                       APPLICATION NUMBER: DE 19932229.5 FILING DATE: 1999-07-09
                                                                                                                                                                                                            APPLICATION NUMBER: DE 19932212.0
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                                                     FILING DATE:
                                                                    APPLICATION NUMBER: DE 19932230.9
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                                                                                                                                           APPLICATION NUMBER: DE 19932228.7
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Zelder, Oskar
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NUMBER: DE 19933005.0
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Pred. No. 7.2e+02;
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                                 ; ORGANISM: Human US-09-949-016-10468
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US-09-949-016-10468
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LENGTH: 630
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                                                                                                     SEQ ID NO 10468
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Best Local
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 600/237,768
                                                                                                                                    PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
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PRIOR FILING DATE:
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FILING DATE: 1999-08-27
APPLICATION NUMBER: DE 19940830.0
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APPLICATION NUMBER: DE 19940833.5
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Pred. No. 7.9e+02;
0; Mismatches 1
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Query Match

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US-09-252-991A-29161
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                                                  RESULT 24
US-09-543-681A-6207
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                ; Sequence 6207, Application US/09543681A
; Patent No. 6605709
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APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                         APPLICANT: GARY BRETON
TITLE OF INVENTION: DUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABIL
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 6306
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GENERAL INFORMATION:
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Matches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 6306, Application US/09543681A Patent No. 6605709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 29161
                                                                                                                                                                                               Best Local Similarity 83.3
Matches 5; Conservative
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                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 675
TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                        ORGANISM: Proteus mirabilis
                                                                                                                                                                                                                                                                                                                          LENGTH: 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                              46
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                                                                                                                                                               1 FSRYAR 6
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5; Conserve
                                                                                                                              FSRYER 51
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83.3%;
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83.3%;
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1; Mismatches 0
                                                                                                                                                                                                                  Score 26;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 27; DB 2;
Pred. No. 8.5e+02;
                                                                                                                                                                                                 Mismatches
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; TOPOLOGY: unl; MOLECULE TYPE: ; MYPOTHETICAL: 1 US-08-497-312-20
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Best Local Similarity
Thes 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILITITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 6207
LENGTH: 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20, Application Patent No. 5712120
                                  Query Match
Best Local Similarity
Matches 5; Conserv
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TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: GARY BRETON
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REGISTRATION NUMBER: 30,549
REFERENCE/DOCKET NUMBER: 2629U
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: OUR SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Proteus mirabilis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 11600
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: CENTRO DE INMUNOLOGIA MOLECULAR
STREET: 215 Y 15, ATABEY PLAYA
                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
LENGTH: 119 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                         TELEPHONE: 801/532-1922
TELEFAX: 801/531-9168
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  Conservative
                                                                                                                                       unknown
                                                                                               : protein
NO
                                                                                                                                                          unknown
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83.9%; Score 26; DB 1; Le
100.0%; Pred. No. 2.4e+02;
. Mismatches 0;
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Pred. No. 2.2e+02;
0; Mismatches 1
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                                        Length 119;
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US-08-497-312-22
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Matches 5; Conserv
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                                                                   GENERAL INFORMAPPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REGISTRATION NUMBER: 30,549
REFERENCE/DOCKET NUMBER: 2629US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 801/531-9168
TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
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APPLICANT:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
              APPLICANT: SIERRA BL ZQUEZ, P.
APPLICANT: TORMO BRAVO, B. R.
TITLE OF INVENTION: Anti-CD6 monoclonal antibodies
                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: p:
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acid
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TELEPHONE: 801/532-1922
TELEFAX: 801/531-9168
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ADDRESSEE: CENTRO DE INMUNOLOGIA MOLECULAR
STREET: 215 Y 15, ATABEY PLAYA
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: uni
TOPOLOGY: unknown
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                                                                                                                       INFORMATION:
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                                                                                                                                                         Application US/08875674A
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                               LOMBARDERO VALLADARES, J.
P REZ RODR GUEZ, R.
SIERRA BL ZQUEZ, P.
TORMO BRAVO, B. R.
                                                                                                        MONTERO
                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  unknown
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                                                                                                                                                                                                                                                                                                                    83.9%; Score 26; DB 1; Long 100.0%; Pred. No. 2.4e+02; tive 0; Mismatches 0;
                                                                                                      CASIMIRO, J. E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method for obtaining modified immunogenicity of murine
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                                                                                                                                                                                                                                             Sequence 3, Application US/08875674A Patent No. 6572857 GENERAL INFORMATION:
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OTHER INFORMATION: Sequence corresponding to the variable region
Patent No. 6572857

OTHER INFORMATION: of the heavy chain of the monoclonal antibody
OTHER INFORMATION: designated as sub-clone for tlA.
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Mice Balb/C INDIVIDUAL ISOLATE: ior tlA TISSUE TYPE: Murine hibridoma IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/CU96/00004
FILING DATE: 118-NOV-1996
ATTORNEY/DAGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
                                                       APPLICANT: LOMBARDERO VALLADARES, J.
APPLICANT: P REZ RODR GUEZ, R.
APPLICANT: SIERRA B.
APPLICANT: TORMO BRAVO, B. R.
TITLE OF INVENTION: Anti-CD6 monoclonal antibodies and their uses
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: Protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: -N Terminal fragment.
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk 3.5' (1.4 MB).
COMPUTER: Compatible PC IBM (80486, 8 M Ram).
OPERATING SYSTEM: Windows 95.
SOFTWARE: Word Perfect 5.0 for Windows 95.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 723-4300
                             NUMBER OF SEQUENCES: 4
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CITY: Scarsdale
STATE: New York
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REFERENCE/DOCKET NUMBER: P-12
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Lackenbach Siegel Marzullo Aronson & Greenspan, P.C.
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                             APPLICANT: Soderlind, Ulf
APPLICANT: Borrebaeck, Carl
TITLE OF INVENTION: A Method For In Vitro Mo
TITLE OF INVENTION: Evolution of Protein Fu
FILE REFERENCE: Mewburn 341711
CURRENT APPLICATION NUMBER: US/10/118,100
CURRENT FILING DATE: 2002-04-08
PRIOR APPLICATION NUMBER: US/09/341,711
PRIOR FILING DATE: 1999-09-21
                                                                                                                                                                                                                                                                           US-10-118-100-51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No. 6572857
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                        Sequence 51, Application US/10118100 Patent No. 6989250
                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                       GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 18-NOV-1996
ATTORNEY AGENT INFORMATION:
NAME: HENRY A. MARZULLO, JR.
REGISTRATION NUMBER: 20,910
REFERENCE/DOCKET NUMBER: P-12
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 723-4300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FRAGMENT TYPE:
ORIGINAL SOURCE:
TISSUE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: PCT/CU96/00004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Word Perfect 5.0 for Win CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,674A
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MEDIUM TYPE: Floppy disk 3.5'' (1.4 MB).
COMPUTER: Compatible PC IBM (80486, 8 M Ram).
OPERATING SYSTEM: Windows 95.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE
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Terminal fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IDENTIFICATION METHOD: By similarity with known sequence. OTHER INFORMATION: Sequence corresponding to the humanized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 119 Amino acid residues.
TYPE: Amino acid.
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APPLICATION NUMBER: PCT/GB98/00219 FILING DATE: 1998-01-26
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SO " SP 2/0 " CHO
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US-09-270-767-36842
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                                                                         US-09-270-767-36842
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PRIOR FILING DATE: 1997-01-24
NUMBER OF SEQ ID NOS: 54
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 51
LENGTH: 130
TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767

CURRENT FILING DATE: 1999-03-17

NUMBER OF SEQ ID NOS: 62517

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 44176

LENGTH: 131
                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Homburger et al.
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION UNMBER: US/09/270,787
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 36842
LENGTH: 165
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                                                                                                                                            TYPE: PRT
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                                                                                         OTHER INFORMATION: Xaa means any amino acid
                                                                                                              FEATURE:
                                                                                                                         ORGANISM: Drosophila melanogaster
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 83.9%; Score 26; DB 2; llarity 100.0%; Pred. No. 3.3e+(Conservative 0; Mismatches
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100.0%; Pred. No. 2.6e+02;
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Pred. No.
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RESULT 34
US-09-549-108-11
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Best Local Similarity
~~~hes 5; Conserva
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; ORGANISM: BR449
US-09-248-528-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 52059
LENGTH: 165
TYPE: PRT
ORGANISM: Drosophila melanogaster
             Sequence 11, Application US/09549108 Patent No. 6214603 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local
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APPLICANT: Padmakumar, Rugmini
APPLICANT: Padmakumar, Rugmini
APPLICANT: Eddmakumar, Rugmini
APPLICANT: Eddmakumar, Rugmini
APPLICANT: Kim, Sang H
TITLE OF INVENTION: Method for Producing Amide Compounds Using a
TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
FILE REFERENCE: MSU 4.1.401
CURRENT APPLICATION NUMBER: US/09/248,528C
CURRENT APPLICATION NUMBER: US/09/248,528C
CURRENT FILING DATE: 1999-02-10
EARLIER FILING DATE: 1999-04-29
RUMBER OF SEQ ID NOS: 18
COMMANDER: DESCRIPTION OF SEQ ID NOS: 18
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                   LENGTH: 229
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Oriel, Patrick J
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Pred. No. 4.6e-
0; Mismatches
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3.3e+02;
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APPLICANT: Oriel, Patrick J
APPLICANT: Padmakumar, Rugmini
APPLICANT: Kim, Sang H
ITITLE OF INVENTION: Method for Producing Amide Compounds Using a l
ITITLE OF INVENTION: Method for Producing Amide Compounds Using a l
ITITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
FILE REFERENCE: MSU 4.1-489
CURRENT APPLICATION NUMBER: US/09/549,111
CURRENT FILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR FILING DATE: 1990-02-10
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
COFFMENDE: Patrick US/
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; ORGANISM: BR449
US-09-549-108-11
                                                                                                                                                                  US-09-549-106-11
                                                                                                                                                                                      RESULT 36
                                                                                                                                                                                                                                                                                                                                                                                                  US-09-549-111-11
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US-09-549-111-11
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CURRENT APPLICATION NUMBER: US/09/549,108
CURRENT FILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 09/248,528
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Pat
SEQ ID NO 11
LENGTH: 229
TYPE: PRT
                                                                                                           Sequence 11, Application US/09549106 Patent No. 6242242 GENERAL INFORMATION:
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Best Local
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Best Local (
               APPLICANT: Oriel, Patrick J
APPLICANT: Padmakumar, Rugmini
APPLICANT: Kim, Sang H
TITLE OF INVENTION: Method for
TITLE OF INVENTION: Hydratase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kim, Sang H
TITLE OF INVENTION: Method for Producing Amide Compounds Using a
TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: BR449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 229
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                                                                                                                                                                                                                                              161 FPRYAR 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    161 FPRYAR 166
                                                                                                                                                                                                                                                                                 1 FSRYAR 6
                                                                                                                                                                                                                                                                                                                     Similarity
5; Conserv
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ON: Method for Producing Amide Compounds Using a ON: Hydratase from a Thermophilic Bacillus MSU 4.1-487
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                                                                                                                                                                                                                                                                                                                     Score 26; DB Pred. No. 4.6e
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Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                 Db -
. 4.6e+02;
1;
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; SOFTWARE: Patent:
; SEQ ID NO 11
; LENGTH: 229
; TYPE: PRT
; ORGANISM: BR449
US-09-550-394-11
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; TYPE: PRT
; ORGANISM: BR449
US-09-549-106-11
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PRIOR TILING DATE: 1998-04-29
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 11
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SEQ ID NO 11
                                                                                                                                                                                                                           Sequence 12718, Application US/09489039A Patent No. 6610836
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Best Local (
SEQ ID NO 12718
LENGTH: 233
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APPLICANT:
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Patent No. 6287828
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                                                     FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
                                                                                                                                            APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/550,394
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 09/248,528
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Padmakumar, Rugmini
APPLICANT: Kim, Sang H
TITLE OP INVENTION: Method for Producing Amide Compounds Using a
TITLE OP INVENTION: Hydratase from a Thermophilic Bacillus
FILE REFERENCE: MSU 4.1-488
CHEBERENCE: MSU 4.1-488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/549,106
CURRENT FILING DATE: 2000-04-13
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                                   NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                     y Match 83.9%;
Local Similarity 83.3%;
hes 5; Conservative
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Pred. No. 4.60
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 26; DB 2;
Pred. No. 4.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
VINUMBER OF SEQ ID NOS: 207012
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6472
LENGTH: 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: PRT
; ORGANISM: Human
US-09-949-016-6472
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                                     ; TYPE: PRT
; ORGANISM: Human
US-09-949-016-9375
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US-09-949-016-9375
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US-09-949-016-6472
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                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS
FILE REFERENCE: CLOO1307
CURRENT APPLICATION UNWERER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
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Patent No. 681233
                                                                                               SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9375
LENGTH: 253
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9375, Application US/09949016 Patent No. 6812339
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  Query Match
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PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Klebsiella pneumoniae
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  83.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26; DB 2;
Pred. No. 5.1e+02;
  Score 26;
  DB
  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 252;
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OF DETECTION AND USES THEREOF
Length 253;
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APPLICANT: Kalman, Sue
APPLICANT: Davis, Ronald
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
FILE REFERENCE: 018941-000411US
CURRENT FILING DATE: 018941-000411US
CURRENT FILING DATE: 2002-03-13
PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR FILING DATE: 1999-11-12
PRIOR FILING DATE: 1999-10-10
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
SOFTWARE: FastSEQ for Windows Version 3.0
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US-09-198-452A-266
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SEQ ID NO 266
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GENERAL INFORMATION:
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Patent No. 6559294
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LENGTH: 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 256, Application US/09438185A Patent No. 6822071
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                                                                                       Query Match
Best Local
                                                                                                                                 TYPE: PRT
ORCANISM: Chlamydia pneumoniae
FEATURE:
OTHER INFORMATION: CPn0254
-09-438-185A-256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Griffais, R.

TITLE OF INVENTION: Chereof and uses thereof, in particular for the diagnos TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnos TITLE OF INVENTION: and treatment of infection FILE REFERENCE: 9710-003-999 CURRENT EPPLICATION NUMBER: US/09/198,452A

CURRENT FILING DATE: 1998-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stephens, Richard APPLICANT: Mitchell, Wayne APPLICANT: Kalman, Sue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 264
TYPE: PRT
ORGANISM: Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 83.9%; So
Local Similarity 100.0%; Figes 5; Conservative 0;
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150 FSRYA 154
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                                                                                       Similarity
                                                                      <u>ب</u>
                                  FSRYA 5
                                                                        Conservative
                                                                83.9%;
                                                                                     Score 26; Pred. No.
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                                                                                     DB 2; Le 5.5e+02;
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FILE REPERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 20485
LENGTH: 283
TYPE: PRT
                                                                                                                                            RESULT 45
US-08-944-483-66
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Best Local Similarity
"~~~hes 5; Conserv?
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US-09-252-991A-20485
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APPLICANT: Marc J. Rubenfield et al.
APPLICANT: MAIC J. RUBENFIELD ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
                                                                                         Sequence 66, Application US/08944483 Patent No. 6232456 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 25159
LENGTH: 276
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence 20485, Application US/09252991A Patent No. 6551795
                                                                                                                                                                                                                                                                                          Matches
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PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788 PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                  APPLICANT:
                                                      APPLICANT:
   APPLICANT:
                                                                                                                                                                                                                                                                                                             Local
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COHEN, MAURICE COLPITTS, TRACEY L. FRIEDMAN, PAULA N. GRANADOS, EDWARD N. KLASS, MICHAEL R.
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 EDWARD N.
CHAEL R.
                                                      TRACEY L.
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Pred. No. 5.7e+02;
1; Mismatches 0
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Pred. No. 5.6e+02;
0; Mismatches 1
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; TYPE: PRT ; ORGANISM: Candida albicans US-09-248-796A-14594
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                                                APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR APPLICATION UNMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NO 14594
LENGTH: 308

TYPE: PET
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                                                                                                                                                                                                                                                                                                                  Sequence 14594, Application US/09248796A Patent No. 6747137
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SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: AND
TITLE OF INVENTION: OF
NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Becker, Cheryl L.
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
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APPLICATION NUMBER: US/08/944,483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
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5; Conserve
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100 Abbott Park Road
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STEVEN D.
NOVEL SERINE PROTEASE REAGENTS
AND METHODS USEFUL FOR DETECTING AND TREATING DISEASES
OF THE PROSTATE
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Pred. No. 6.1e
0; Mismatches
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CURRENT FILING DATE: 2000-01-27

PRIOR APPLICATION NUMBER: US 60/117,747

PRIOR FILING DATE: 1999-01-29

NUMBER OF SEQ ID NOS: 14342

SEQ ID NO 14005

LENGTH: 327

TYPE: PRT

ORGANISM: Klebsiella pneumoniae
US-09-489-039A-14005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Darrow, Andrew
APPLICANT: Qi, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 12
LENGTH: 319
TYPE: PRT
TYPENTY Artificial Sequence
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US-09-489-039A-14005
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US-09-386-642-12
; Sequence 12, Application US/09386642
; Patent No. 6420157
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Best Local
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Best Local
                                                                 Best Local Similarity
Matches 5; Conserv
                                                                                                   Query Match
                                                                                                                                                                                                                                                                             FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
                                                                                                                                                                                                                                                                                                                             APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Fusion OTHER INFORMATION: with homo sapien serine protease catalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
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251 WSRYAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      144 FSRYIR 149
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5; Conserv
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Pred. No. 6.2e+02;
                                                                     Score 26; DB 2; Le
Pred. No. 6.6e+02;
1; Mismatches 0;
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Pred. No. 6.5e+02;
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                                                                                                     Length 327;
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RESULT 49 US-09-386-642-11 ; Sequence 11, Application US/09386642

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APPLICANT: Darrow, Andrew
APPLICANT: Darrow, Andrew
APPLICANT: Canton, Patricia
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION UNMBER: US/09/386,642
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn Ver. 2.0
SOFTWARE: PatentIn Ver. 2.0
SOFTWARE: PRI
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Fusion gene
OTHER INFORMATION: with homo sapien serine protease catalytic domain
US-09-386-642-11
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APPLICANT: Stephens, Richard
APPLICANT: Mitchell, Wayne
APPLICANT: Mitchell, Wayne
APPLICANT: Kalman, Sue
APPLICANT: Kalman, Sue
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
FILE REFERENCE: 0.18941-000411US
CURRENT APPLICATION NUMBER: US/09/438,185A
CURRENT FILING DATE: 2002-03-13
FRIOR APPLICATION NUMBER: US 60/108,279
PRIOR APPLICATION NUMBER: US 60/108,279
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR APPLICATION NUMBER: US 60/128,606
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 1074
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 960
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Search completed: August 29, 2006, 06:17:52 Job time : 14.3366 secs
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                                                                                                                                                                                                      Query Match 83.9%; Score 26; DB 2; Length 337; Best Local Similarity 83.3%; Pred. No. 6.8e+02; Matches 5; Conservative 0; Mismatches 1; Indels
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Sequence 542	Sequence 23, Sequence 23, Sequence 24, Sequence 542, Seque
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and Methods of Use Length 6; Indels 0; Gaps 0;	Sequence 542, App Sequence 542

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RESULT 4
US-10-541-343-6
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; Publication No. US20060052299A1
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Publication No. US20060052299A1
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APPLICANT: Ko, Pui-Yuk Dorothy
TITLE OF INVENTION: MNTF Peptides and Compositions and Methods
FILE REFERENCE: 2001-103US
CURRENT APPLICATION NUMBER: US/10/541,343
CURRENT FILING DATE: 2005-07-05
PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR FILING DATE: 2004-01-21
PRIOR FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 60/441,722
PRIOR FILING DATE: 2003-01-21
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PRIOR FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 60/441,722
PRIOR FILING DATE: 2003-01-21
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                  OTHER INFORMATION: Synthetic peptide -10-541-343-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Chau, Raymond Ming Wah
APPLICANT: Ko, Pui-Yuk Dorothy
TITLE OF INVENTION: MOTER Peptides and Compositions and Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.3
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CURRENT APPLICATION NUMBER: US/10/541,343
CURRENT FILING DATE: 2005-07-05
                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial
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OTHER INFORMATION: Synthetic peptide
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CURRENT APPLICATION NUMBER: US/10/541,343
CURRENT FILING DATE: 2005-07-05
PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR FILING DATE: 2004-01-21
PRIOR FILING DATE: 2003-01-21
PRIOR FILING DATE: 2003-01-21
VUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.3
SEQ ID NO 6
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                                                                        Sequence 4, Application US/09989481 Publication No. US20020086831A1 GENERAL INFORMATION:
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APPLICANT: Chau, Raymond Ming Wah
APPLICANT: KO, Pui-Yuk Dorothy
TITLE OF INVENTION: MNTF Peptides and Compositions and Methods of Use
FILE REFERENCE: 2001-103US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.3 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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Publication No. US20060052299A1
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APPLICANT: Ko, Pui-Yuk Dorothy
TITLE OF INVENTION: MNTF Peptides and Compositions and Methods of Use
FILE REFERENCE: 2001-103US
CURRENT APPLICATION NUMBER: US/10/541,343
CURRENT FILING DATE: 2005-07-05
PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR PILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 60/441,722
PRIOR FILING DATE: 2003-01-21
APPLICANT: Chau, Raymond M.W.
TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-4
CURRENT APPLICATION NUMBER: US/09/989,481
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TYPE: PRT
ORGANISM: Artificial
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Synthetic peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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Pred. No.
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US-10-858-286-4
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US-10-858-545-4
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                                                                                                            Sequence 4, Application US/10858286
Publication No. US20040242486A1
GENERAL INFORMATION:
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Publication No. US20040224894A1
GENERAL INFORMATION:
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Matches 6; Conserv
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PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR FILING DATE: 1996-09-27
NUMBER OF SEQ ID NOS: 5
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APPLICANT: Chau, Raymond M.W.
TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-3
CURRENT APPLICATION NUMBER: US/10/858,286
CURRENT FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/09/592,018
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PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR EILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
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CURRENT APPLICATION NUMBER: US/10/858,545
CURRENT FILING DATE: 2004-06-01
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
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TYPE: PRT
ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
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Pred. No.
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CURRENT APPLICATION NUMBER: US/10/858,543
CURRENT FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/09/592,018
PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR APPLICATION NUMBER: US 08/9528662
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
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PRIOR FILLING DATE: 1997-09-12
PRIOR PELICATION NUMBER: US 08/751225
PRIOR PILLING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR APPLICATION NUMBER: US 60/026792
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.0
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LENGTH: 33
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Publication No. US20040266691A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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FILE REFERENCE: 12592-3
CURRENT APPLICATION NUMBER: US/10/858,144
CURRENT FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/09/592,018
PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
                                                                                                                                                APPLICANT: Chau, Raymond M.W. TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors FILE REFERENCE: 12592-3
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
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NUMBER OF SEQ ID NOS:
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TYPE: PRT
ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
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GENERAL INFORMATION:

APPLICANT: Abad, Mark S. et al.

FILE REFERENCE: 38-21(53452)B

CURRENT APPLICATION NUMBER: US/11/188,298

CURRENT FILING DATE: 2005-07-22

PRIOR APPLICATION NUMBER: 60/592,978

PRIOR FILING DATE: 2004-07-31

NUMBER: OF SEQ ID NOS: 22589

SEQ ID NO 11746

LENGTH: 125

TYPE: PRT
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US-10-541-343-1
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US-11-188-298-11746
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US-11-188-298-11746
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Best Local o
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PRIOR APPLICATION NUMBER: US 60/026792
PRIOR FILING DATE: 1996-09-27
NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 60/441,722
PRIOR FILING DATE: 2003-01-21
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Chau, Raymond Ming Wah
APPLICANT: Ko, Pui-Yuk Dorothy
TITLE OF INVENTION: WATE Peptides and Compositions and Methods of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.0
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CURRENT FILING DATE: 2005-07-05
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TYPE: PRT
ORGANISM: Artificial
FEATURE:
          ORGANISM: Desulfitobacterium hafniense
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                                                          US-10-289-762-977
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US-10-289-762-977
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PRIOR FILING DATE: 2000-07-11
PRIOR PPLICATION NUMBER: 0017983.8
PRIOR APPLICATION NUMBER: 0017983.8
PRIOR APPLICATION NUMBER: 0019368.0
PRIOR FILING DATE: 2000-08-07
PRIOR PRIOR APPLICATION NUMBER: 0020440.4
PRIOR PRIOR DATE: 2000-08-18
PRIOR APPLICATION NUMBER: 0027583.9
PRIOR APPLICATION NUMBER: 0027549.5
PRIOR APPLICATION NUMBER: 0027549.5
                                                                                                                                                                                                                            Sequence 977, Application US/10289762
Publication No. US20040006218A1
GENERRAL INFORMATION:
APPLICANT: Griffais, R.
APPLICANT: Griffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preventing of INVENTION: and treatment of infection
                                                                                                                               FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/10/289,762
CURRENT FILING DATE: 2003-03-27
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 977
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SEQ ID NO 107
LENGTH: 357
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Best Local Similarity
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CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: 001633.4
PRIOR FILING DATE: 2000-07-03
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TITLE OF INVENTION: IMMUNISATION AGAINST CHLAMYDIA PNEUMONIAE
FILE REFERENCE: P025035WO
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                                                                          TYPE: PRT ORGANISM: Chlamydia pneumoniae
                                                                                                                 LENGTH: 357
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Pred. No. 1.3e+02;
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Pred. No.
  Score 31; DB 4;
Pred. No. 1.3e+02;
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Mismatches
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RESULT 16
US-10-437-963-176098
; Sequence 176098, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
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Best Local Similarity
Thes 6; Conserv
                                                                                                                                                                                                                                                                    ; ORGANISM: Chlamydia pneumoniae US-10-282-122A-54956
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                                                                                                                                                                                                                                                                                              Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 78614
SOPTWARE: PatentIn version 3.1
SEQ ID NO 54956
LENGTH: 357
TYPE: PRT
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APPLICANT:
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APPLICANT: Zamuc
APPLICANT: Malor
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PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/230,335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Identification of Essential Genes in Microorganisms FILE REFERENCE: ELITRA.034A CURRENT APPLICATION NUMBER: US/10/282,122A CURRENT FILING DATE: 2003-02-20
APPLICANT:
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APPLICATION NUMBER: 60/267,636
FILING DATE: 2001-02-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 2000-09-06
APPLICATION NUMBER: 60/230,347
FILING DATE: 2000-09-09
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 60/269,308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/253,625 FILING DATE: 2000-11-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/257,931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 2000-10-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/242,578
                                                                                                                                          135 FSRYAR 140
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La Rosa, Thomas
Kovalic, David
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Malone, Cheryl
Haselbeck, Robert
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Zyskind, Judith
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Carr, Grant
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Pred. No. 1.3e+02;
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US-10-369-493-23258
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Best Local (
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TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN I
TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
FILE REFERENCE: 38-10 (52052)B
CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US-10/369,493
PRIOR FILING DATE: 2002-02-21
NUMBER OF SEQ ID NOS: 47374
SEQ ID NO 23258
LENGTH: 463
TYPE: PRT
ORGANISM: Bacillus subtilis
US-10-369-493-23258
Sequence 1759, Application US/11087099

Publication No. US20060041961A1

GENERAL INFORMATION:

APPLICANT: Abad, Mark S. et al.

TITLE OF INVENTION: Genes and Uses for Plant

FILE REFERENCE: 38-21(53450)B EP

CURRENT APPLICATION NUMBER: US/11/087,099

CURRENT APPLICATION NUMBER: US/11/087,099

NUMBER OF SEQ ID NOS: 12464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23258, Application US/10369493
Publication No. US20030233675A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Cao, Yongwei
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Barry S.
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6; Conserva
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Wu, Wei
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Pred. No. 1.5e+02;
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                                                                                    for Plant Improvement
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US-11-188-298-12707
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; TYPE: PRT
; ORGANISM: Bacillus subtilis subsp. subtilis str. 168
US-11-087-099-1759
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US-11-188-298-12707
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILLING DATE: 2000-03-21
PRIOR PELLING DATE: 2000-03-21
PRIOR PELLING DATE: 2000-05-23
PRIOR PELLING DATE: 2000-05-26
PRIOR PELLING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILLING DATE: 2000-10-23
PRIOR PELLING DATE: 2000-11-27
PRIOR PELLING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR PELLING DATE: 2000-11-27
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PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 12707
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APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21 (53452) B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
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Best Local Similarity 100.0%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                      APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITER.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Haselbeck, Robert APPLICANT: Ohlsen, Kari L. APPLICANT: Zyskind, Judith W.
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APPLICANT:
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Trawick, John D.
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APPLICANT: Hyber, Inc
APPLICANT: Hyber, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POI
FILE REFERENCE: 790C1P3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
NUMBER OF SEQ ID NOS: 60736
SOPTWARE: CUSTOM
SEQ ID NO 54803
LENGTH: 1225
TYPE: PRT
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US-10-369-493-23518
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US-10-369-493-23518
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US-09-815-242-10270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES FILE REFERENCE: 38-10 (52052)B CURRENT APPLICATION NUMBER: US/10/369,493 CURRENT FILING DATE: 2003-02-28 PRIOR APPLICATION NUMBER: US 60/360,039 PRIOR FILING DATE: 2002-02-21 NUMBER OF SEQ ID NOS: 47374 SEQ ID NO 23518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23518, Application US/10369493 Publication No. US20030233675A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     Sequence 54803, Application US/10450763 Publication No. US20050196754A1 GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10270
LENGTH: 474
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Matches 6; Conserv
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Best Local (
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APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Goldman, Barry S.
APPLICANT: Chen, Xianfeng
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 100.0%; Score 31; DB 4; Similarity 100.0%; Pred. No. 1.7e+02; 6; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 1.7e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                 POLYPEPTIDES
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FILTE OF INVESSION: NOTE: TITLE OF INVESSION: NUMBER: US/10/450,763; CURRENT APPLICATION NUMBER: US/10/450,763; CURRENT FILING DATE: 2003-06-11; PRIOR APPLICATION NUMBER: PCT/US01/08631; PRIOR FILING DATE: 2001-03-30; PRIOR PRIOR DATE: 2001-03-30; PRIOR PRIOR DATE: 2000-03-31; PRIOR PRIOR DATE: 2000-03-31; PRIOR PRIOR DATE: 2000-03-31; PRIOR PRIOR DATE: 2000-08-23; NUMBER: OPSEQ ID NOS: 60736; SOFTWARE: Custom SEQ ID NO 55178
RESULT 24
US-10-424-599-186222
; Sequence 186222, Application US/10424599
; Publication No. US20040031072A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-10-450-763-55178; Sequence 55178, Application US/10450763; Publication No. US20050196754A1
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Best Local Similarity
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TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sap
FEATURE:
NAME/KEY: DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: DOMAIN
LOCATION: (332)..(380)
LOCATION: Tetrahydrofolate dehydrogenase/cyclohydrolase proteins domain
OTHER INFORMATION: identified by eMATRIX, accession number BL00768C, p-value=1.000e-
OTHER INFORMATION: 40, raw score of 25.86
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OTHER INFORMATION: Tetral
OTHER INFORMATION: PFam,
OTHER INFORMATION: 460.9
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                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Tetrahydrofolate dehydrogenase/cycloh domain identified by OTHER INFORMATION: PFam, accession name THF_DHG_CYH, E-value=2.9e-134, PFam score OTHER INFORMATION: 459.5
                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: DOMAIN
LOCATION: (267)..(471)
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LOCATION: (7)..(
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Pred. No. 4.4e+02;
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Pred. No.
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ches 0;
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APPLICANT: Covalic David K
APPLICANT: Cabou Yihua
APPLICANT: Cabou Yihua
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285884
SEQ ID NO 146582
LENGTH: 79
Type----
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APPLICANT: La Rosa Thomas J

APPLICANT: Kovalic David K

APPLICANT: Zhou Yihua

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Sy Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement

FILE REFERENCE: 38-21(5323)B

CURRENT APPLICATION NUMBER: US/10/424,599

CURRENT APPLICATION NUMBER: US/10/424,599

RUMBER OF SEQ ID NOS: 285684
                                                                     Sequence 265564, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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LENGTH: 67
TYPE: PRT
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Best Local Similarity
Matches 5; Conserv
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                                      APPLICANT: La Rosa Thomas APPLICANT: Kovalic David
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ORGANISM: Glycine max
FEATURE:
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LOCATION: (1)..(67)
OTHER INFORMATION: unsure at
FEATURE:
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Kovalic David K
Zhou Yihua
Cao Yongwei
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Pred. No. 1.4e+02;
1; Mismatches 0
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Pred. No. 1.1e+02;
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APPLICANT:

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US-10-424-599-265564
                                                                                                      ; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (85)
; OTHER INFORWATION: Xaa =
US-10-183-687-178
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Matches
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APPLICANT: Allen, Willia
APPLICANT: Cahoon, Rebec
APPLICANT: Epelbaum, Said
APPLICANT: Famodu, Omoli
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LENGTH: 85
                                           Matches
                                                          Query Match
Best Local (
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APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                           FEATURE:
NAME/KEY: UNSURE
LOCATION: (80)
OTHER INFORMATION: >
                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/301,913
PRIOR FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Microsoft Office 97
IEQ ID NO 178
                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Tarczynski, Mitchell C.
TITLE OF INVENTION: Alteration Of Oil Traits
FILE REFERENCE: BB1458 US NA
CURRENT APPLICATION NUMBER: US/10/183,687
CURRENT FILING DATE: 2002-06-27
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ORGANISM: Glycine max
FEATURE:
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5; Conserv
               1 FSRYAR 6
                                          Similarity
5; Conserv
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Shen, Bo
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Harvell, Leslie T.
Jones, Todd
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Oliveira, Igor Cunha
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Klein, Ted
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o. US20030204870A1
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                                                          90.3%;
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                                           1; Mismatches
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Pred. No. 1.5e+02;
1; Mismatches (
                                                          Score 28; I
Pred. No. 1.
                                                                                                                     acid
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1.5e+02;
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                                           0; Indels
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                                                                      Length 90;
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RESULT 30
US-10-767-701-43382
US-10-767-701-43382, Application US/10767701
; Sequence 43382, Application US/20767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
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US-10-767-701-51064
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Best Local
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FSKYAR
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APPLICANT: Zhou, Yihua

APPLICANT: Zhou, Yihua

APPLICANT: Zhou, Yihua

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21 (53222) B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 219020

LENGTH: 91

TYPE: PT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Clone ID: MRT4577_131339C.1.pep US-10-425-115-219020
                                                                            ; OTHER INFORMATION: Clone ID: LIB3480-055-P1-K1-H11.pep US-10-767-701-51064
                                                                                                                            GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof For Plant Improvement
FILE REFERENCE: 38-21(53535)B
CURRENT FILIG DATE: 2004-01-29
NUMBER OF SEQ ID NOS: 63128
SEQ ID NO 51064
LENGTH: 123
TYPE: PRT
ORGANISM: Sorghum bicolor
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51064, Application US/10767701 Publication No. US20040172684A1
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                    90.3%;
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Score 28; DB 4;
Pred. No. 2.1e+02;
1; Mismatches (
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                                      Length 123;
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                                                                                                                                                                                                                                                                                                        APPLICANT: Tarczyński, Mitchell C.
TITLE OF INVENTION: Alteration Of Oil Traits In Plants
FILE REFERENCE: BB1458 US NA
CURRENT APPLICATION NUMBER: US/10/183,687
CURRENT FILING DATE: 2002-06-27
PRIOR APPLICATION NUMBER: 60/301,913
PRIOR FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Microsoft Office 97
SEQ ID NO 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 43382
LENGTH: 124
TYPE: PRT
ORGANISM: Sorghum bicolor
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NAME/KEY: UNSURE
LOCATION: (102)
OTHER INFORMATION: :
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                                                                                                                                                                                                                                          LENGTH: 132
TYPE: PRT
ORGANISM: Oryza sativa
FEATURE:
                                                                                                                                                                                            NAME/KEY: UNSURE LOCATION: (91)
OTHER INFORMATION:
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 OTHER INFORMATION:
                                                               OTHER INFORMATION:
                                 NAME/KEY: UNSURE
                OCATION:
                                                                               AME/KEY: UNSURE
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Local Similarity 83.3%;
les 5; Conservation
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Cao, Yongwei
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Oliveira, Igor Cunha
Sakai, Hajime
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Harvell, Leslie T.
Jones, Todd
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Klein, Ted
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Xaa
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Pred. No. 2.1e+02;
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; NAME/KEY: UNSURE
; LOCATION: (122)
; OTHER INFORMATION: X
FEATURE:
; PEATURE:
; NAME/KEY: UNSURE
; LOCATION: (125)
; OTHER INFORMATION: X
US-10-183-687-174
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Best Local Similarity
~~~hes 5; Conserva
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US-10-501-282-480
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US-10-501-282-478
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SEQ ID NO 478
LENGTH: 145
TYPE: PRT
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CURRENT APPLICATION NUMBER: US/10/501,282 CURRENT FILING DATE: 2004-07-09 PRIOR APPLICATION NUMBER: 60/333,777 PRIOR FILING DATE: 2001-11-29 PRIOR APPLICATION NUMBER: 60/426,742 PRIOR FILING DATE: 2002-11-18
                                                                                                                         TITLE OF INVENTION: ALLOIOCOCCUS OTITIDIS OPEN READING TITLE OF INVENTION: POLYPEPTIDE ANTIGENS, IMMUNOGENIC FILE REFERENCE: AM100780 L2
                                                                                                                                                                                    APPLICANT: MCMICHAEL, JOHN CALHOUN
APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
APPLICANT: FLETCHER, LEAH DIANE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/333,777
PRIOR TILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 60/426,742
PRIOR FILING DATE: 2002-11-18
PRIOR PELICATION NUMBER: PCT/US02/36123
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 6653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: AM100780 L2
CURRENT APPLICATION NUMBER: US/10/501,282
CURRENT FILING DATE: 2004-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
APPLICANT: RUSSELL, DAVID PARRISH
APPLICANT: FLETCHER, LEAH DIANE
TITLE OF INVENTION: ALLOIOCOCCUS OTITIDIS OPEN READING FRAMES (ORFS) ENCODING
TITLE OF INVENTION: POLYPEPTIDE ANTIGENS, IMMUNOGENIC COMPOSITIONS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MCMICHAEL, JOHN CALHOUN
APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
APPLICANT: FLETCHER, LEAH DIANE
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Pred. No. 2.5e
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2.5e+02;
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                                                                                                                                              FRAMES (ORFS) ENCODING COMPOSITIONS AND USES
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RESULT 34
US-09-984-245-156
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PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 6653
SOFTWARE: PATENTIN VETSION 3.2
SEQ ID NO 480
LENGTH: 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 156,
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Best Local
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PRIOR
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PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: PCT/US98/09
PRIOR FILING DATE: 1998-03-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/984,245
CURRENT FILING DATE: 2001-10-29
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TITLE OF INVENTION: 87 Human Secreted Proteins
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FILING DATE: 1998-03-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 1997-03-21
APPLICATION NUMBER: US 60/041,276
                                                                                                                                                                                                           FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,352
FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 1997-05-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/048,094
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APPLICATION NUMBER: US 60/048,069
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,096
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PRIOR APPLICATION NUMBER: US 60/054,804
PRIOR FILING DATE: 1997-08-05
PRIOR APPLICATION NUMBER: US 60/056,370
PRIOR FILING DATE: 1997-08-19
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PRIOR FILING DATE: 1997-10-02
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PRIOR APPLICATION NUMBER: US 09/154,707
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: PCT/US98/05311
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CURRENT FILING DATE: 2001-10-01
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hes 5; Conserv
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                                            APPLICATION NUMBER: US (FILING DATE: 1997-05-30
                                                                                                                           APPLICATION NUMBER: US 60/048,186
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APPLICATION NUMBER: US
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Pred. No. 2.9e+02
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; TYPE: PRT ; ORGANISM: Homo sapiens US-09-966-262-156
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US-09-983-966-156
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 156
LENGTH: 174
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PRIOR FILING DATE: 1998-03-19
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CURRENT FILING DATE: 2001-10-29
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TITLE OF INVENTION: 87 Human Secreted Proteins
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FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,160
FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,351
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,352
FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,186
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US
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APPLICATION NUMBER: US 60/041,276
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FILING DATE: 1997-08-19
APPLICATION NUMBER: US 60/060,862
FILING DATE: 1997-10-02
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APPLICATION NUMBER: US 60/054,804
FILING DATE: 1997-08-05
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APPLICATION NUMBER: US 60/048,099
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10MBbx. 1997-05-30 US 60/048,069
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Pred. No.
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PRIOR APPLICATION NUMBER: US/09/966,262
PRIOR FILING DATE: 1001-10-01
PRIOR PELLING DATE: 1998-09-17
PRIOR FILING DATE: 1998-09-17
PRIOR FILING DATE: 1998-09-17
PRIOR FILING DATE: 1998-03-19
PRIOR PELLING DATE: 1998-03-19
PRIOR PELLING DATE: 1997-03-21
PRIOR PELLING DATE: 1997-03-30
PRIOR PELLING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/048, 350
PRIOR APPLICATION NUMBER: US 60/048, 350
PRIOR PELLING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/048, 135
PRIOR PELLING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/048, 135
PRIOR PELLING DATE: 1997-05-30
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LENGTH: 174
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CURRENT FILING DATE: 2002-01-31
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FILE REFERENCE: PZ004P1
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NUMBER OF SEQ ID NOS: 343
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ORGANISM: Homo
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APPLICATION NUMBER: US 60/056,370
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APPLICATION NUMBER: US 60/048,096
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o. US20030018180A1
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83.3%;
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Pred. No. 2.9e+02;
Pred. No. 2.9e+02;
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APPLICATION NUMBER: US 60/048,186 FILING DATE: 1997-05-30

APPLICATION NUMBER: US 60/048,352 FILING DATE: 1997-05-30

FILING DATE: 1997-05.

APPLICATION NUMBER: US 60/048,099

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                                                                                                                                     US-10-143-090-156
                                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 156
                                                                                                                                                                                                                                                                                                                                                                       Sequence 156, Application US/10143090
Publication No. US20030069406A1
GENERAL INFORMATION:
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Best Local S
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                                                                                 Query Match
Best Local
                                                                  Matches
                                                                                                                                                                                                                                                                                APPLICANT: Young et al.
TITLE OF INVENTION: 87 Human Secreted Proteins
FILE REPERENCE: PZOO4P1
CURRENT APPLICATION NUMBER: US/10/143,090
CURRENT FILING DATE: 2002-05-13
                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/154,707
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 343
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TYPE: PRT
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ORGANISM: Homo sapiens
                                                                                                                                                                                    ENGTH: 174
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APPLICATION NUMBER: US 60/060,862
FILING DATE: 1997-10-02
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APPLICATION NUMBER: US 60/048,355
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,095
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                                                                 Similarity 5; Conserv
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Similarity 83.3%;
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FARYAR 74
                                                                  Conservative
                                                                                   90.3%;
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Pred. No. 2.9e+02
                                                                 Score 28; DB 4;
Pred. No. 2.9e+02;
1; Mismatches 0
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RESULT 40
US-10-437-963-197182
; Sequence 197182, Application US/10437963
; Publication No. US20040123343A1
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US-10-960-251-156
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US-10-960-251-156
                                                                            APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT PILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 197182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 156, Appropriate Publication No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/154,707
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: PCT/US98/05311
PRIOR FILING DATE: 1998-03-19
PRIOR APPLICATION NUMBER: 60/041,277
PRIOR FILING DATE: 1997-03-21
PRIOR APPLICATION NUMBER: 60/042,344
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CURRENT FILING DATE: 2004-10-08
PRIOR APPLICATION NUMBER: US 10/143,090
PRIOR FILING DATE: 2002-05-13
                                                                                                                                                                                                                                                   APPLICANT:
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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TITLE OF INVENTION: 87 Human Secreted Proteins
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PRIOR FILING DATE: 1997-03-21
PRIOR APPLICATION NUMBER: 60/041,281
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                                        TYPE: PRT
                    ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 174
FEATURE:
                                                              ENGTH: 179
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Wu, Wei
                                                                                                                                                                                                                                                 Barbazuk, Brad
                                                                                                                                                                                                                                                                       Boukharov, Andrey A.
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5. US20050089911A1
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Pred. No. 2.9e+02;
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                                                                                                                         ; OTHER INFORMATION: Clone ID: MRT4577_4982C.1.pep US-10-425-115-314441
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US-10-425-115-314441
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US-11-096-568A-9600
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APPLICANT: Alexandrov, Nickolai et al.

TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides

TITLE OF INVENTION: Therby

FILE REFERENCE: 2750-1592PUS2

CURRENT FILING DATE: 2005-04-01

NUMBER OF SEQ ID NOS: 34471

SEQ ID NO 9600
                                                                                                                                                                                   APPLICANT: La ROSa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 314441
LENGTH: 234
TYPE: PRT
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Matches
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Publication No. US20040214272A1
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LOCATION: (1)..(225)
OTHER INFORMATION: Ceres Seq. ID no. 12614812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Triticum aestivum
FEATURE:
                                                                                                                                                       ORGANISM: Zea mays FEATURE:
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                                                         Local Similarity
ses 5; Conserv
 153
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                             1 FSRYAR 6
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FSKYAR 158
                                                            Conservative
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; Pred. No. 3e+0
1; Mismatches
                                                            1; Mismatches
                                                                          Score 28;
Pred. No.
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Pred. No. 3.8e+02;
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                                                                          DB 4;
3.9e+02;
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3e+02;
                                                       0;
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                                                                                          Length 234;
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                                                            Indels
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APPLICANT: Caovalic David K
APPLICANT: Cabo Yihua

APPLICANT: Cao Yongwei

TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 254640
LENGTH: 239
RESULT 45
US-10-183-687-160
(S-10-183-687-160)
; Sequence 160, Application US/10183687
; Publication No. US20030204870A1
; GENERAL INFORMATION:
; APPLICANT: Allen, Steve
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                                                                                                                                                                                                                                                                                            ; TYPE: PRT ; ORGANISM: Arabidopsis thaliana US-10-987-454-122
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US-10-987-454-122
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                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 251
SOFTWARE: PatentIn version 3.3
SEQ ID NO 122
LENGTH: 243
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CURRENT APPLICATION NUMBER: US/10/987,454
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: US 60/519,606
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: US 60/579,733
PRIOR APPLICATION NUMBER: US 60/579,733
PRIOR FILING DATE: 2004-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF APOLIPOPROTEINS IN TRANSGENIC
TITLE OF INVENTION: PLANTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Glycine max
FEATURE:
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5; Conserva
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Pred. No. 4.1e+02;
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Pred. No. 4e+02;
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APPLICANT:

Allen, William B.

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APPLICANT: Shen, Bo
APPLICANT: Tarczynski, Mitchell C.
FITLE OF INVENTION: Alteration Of Oil Traits In Plants
FILE REFERENCE: BB1458 US NA
CURRENT APPLICATION NUMBER: US/10/183,687
CURRENT FILING DATE: 2002-06-27
PRIOR APPLICATION NUMBER: 60/301,913
PRIOR FILING DATE: 2001-06-29
NUMBER OF SEQ ID NOS: 532
SOFTWARE: Microsoft Office 97
SEQ ID NO 160
LENGTH: 244
TYPE: PRT
ORGANISM: Zea mays
US-10-183-687-160
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US-10-425-114-54567
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RESULT 47
US-10-437-963-133297
; Sequence 133297, Application US/10437963
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                                                                                                                                                                                                                                                                                                                     APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E
APPLICANT: Screen, Steven E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 54567
LENGTH: 256
                                                                                                                                                        Query Match
Best Local Similarity
Matches 5; Conserv
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Best Local Similarity 83.3%;
Matches 5; Conservative
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                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Glycine max FEATURE:
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                                                                                         174 FSKYAR 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 FSRYAR 6
                                                                                                                             1 FSRYAR 6
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Oliveira, Igor Cunha
Sakai, Hajime
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Klein, Ted
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Harvell, Leslie T.
Jones, Todd
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Pred. No. 4.3e+02;
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; Publication AV.; GENERAL INFORMATION: APPLICANT: La Rosa, Thomas J.; APPLICANT: Kovalic, David K.; APPLICANT: Zhou, Yihua; APPLICANT: Zhou, Yongwei
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RESULT 49
US-10-987-454-128
                                                                                                                                                                                                                      ; ORGANISM: Hordeum vulgare US-10-987-454-127
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Sequence 127, Application US/10987454
Publication No. US20050172359A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.3
SEQ ID NO 127
LENGTH: 301
TYPE: PRT
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TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 133297
LENGTH: 285
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Best Local Similarity
Matches 5; Conserv
                                                                                                                                                Matches
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                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/987,454
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: US 60/519,606
PRIOR FILING DATE: 2003-11-14
PRIOR APPLICATION NUMBER: US 60/579,733
PRIOR FILING DATE: 2004-06-16
NUMBER OF SEQ ID NOS: 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Reid, Alexandra APPLICANT: Moloney, Maurice TITLE OF INVENTION: METHODS FILTE OF INVENTION: PLANTS FILE REFERENCE: 9369-311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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ORGANISM: Oryza sativa
FEATURE:
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                                                                                                                                                                Local
                                                                         220 FSKYAR 225
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                                                                                                                                              Similarity 5; Conserv
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Barbazuk, Brad
                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 28; DB 4;
Pred. No. 4.8e+02;
1; Mismatches 0
                                                                                                                                                                  Score 28; DB 5; Pred. No. 5e+02;
                                                                                                                                                Mismatches
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Sequence 128, Application US/10987454
Publication No. US20050172359A1
GENERAL INFORMATION:
APPLICANT: Reid, Alexandra
APPLICANT: Moloney, Maurice

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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF APOLIPOPROTEINS IN TRANSGENIC TITLE OF INVENTION: PLANTS
FILE REFERENCE: 9369-311
CURRENT APPLICATION NUMBER: US/10/987,454
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: US 60/519,606
PRIOR FILING DATE: 2003-11-14
PRIOR FILING DATE: 2003-11-14
PRIOR FILING DATE: 2004-06-16
PRIOR FILING DATE: 2004-06-16
PRIOR FILING DATE: 2004-06-16
VARIANCE: PATOR FILING DATE: 2004-06-16
VARIANCE: PATOR FILING DATE: 2004-06-16
VARIANCE: PATOR OF SEQ ID NOS: 251
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 128
LENGTH: 301
TYPE: PRT
CRGANISM: Hordeum vulgare
US-10-987-454-128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 37522, Application US/10450763

Publication No. US20050196754A1

GENERAL INFORMATION:
APPLICANT: Hyseq, Inc
TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
FILE REFERENCE: 790C1P3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: 09/540,167
PRIOR FILING DATE: 2000-08-23
NUMBER OF SEQ ID NOS: 60736
SOFTWARE: CUSTOM
SEQ ID NO 37522
LENGTH: 301
TYPE: PRI
TY
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FEATURE:
FEATURE
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Search completed: August 29, 2006, 10:18:53
Job time: 75.5941 secs
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US-10-450-763-37522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 90.3%;
Best Local Similarity 83.3%;
Matches 5; Conservative
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Local Similarity 83.3%;
nes 5; Conservative
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90 FARYAR 95
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Pred. No. 5e+02;
1; Mismatches
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S-11-056-355B-85574

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US-11-105-233-53

Sequence 53, Application US/11105233

Publication No. US20060134653A1

GENERAL INFORMATION:
APPLICANT: Thiagalingam et al
TITLE OF INVENTION: Differential Expression of Ger
TITLE OF INVENTION: Tumors
FILE REFERENCE: 1657/2001
CURRENT APPLICATION NUMBER: US/11/105,233
CURRENT FILING DATE: 2005-04-13
NUMBER OF SEQ ID NOS: 202
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 53
LENGTH: 174
                                                                                                                         ; LENGTH: 174
; TYPE: PRT
; ORGANISM: Homo &
US-11-105-233-53
RESULT 2
US-11-056-355B-68811
; Sequence 68811, Application US/11056355B
                                                                                     Query Match
Best Local S
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US-11-30-40-170-8
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US-11-23-49-902-32618
US-11-056-3558-3697-3462
US-11-056-3558-107948
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US-11-056-3558-15736
US-11-056-3558-157736
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                                                                                     score 28; DB; Pred. No. 28; 1; Mismatches
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                                                                                                       Length 174;
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; TYPE: prt
; CRGANISM: Arabidopsis thaliana
; PERTURE:
; NAME/KEY: peptide
; LCCATION: (1)..(240)
; OTHER INFORMATION: Ceres Seq. ID no. 13619980
US-11-056-355B-106575
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US-11-056-355B-106575
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                                                                                                                                    RESULT 4
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Best Local Similarity 83...
Thes 5; Conservative
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PRIOR APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 68811
                                                          Sequence 117814, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 106575
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590FUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
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                       APPLICANT: Brover, Vyacheslav APPLICANT: Alexandrov, Nicko:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: prt
ORGANISM: Triticum aestivum
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(225)
OTHER INFORMATION: Ceres Seq. ID no.
PPLICANT: Alexandrov, Nickolai
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                                                                                                                                                                                         156 FSKYAR 161
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APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
FTILE OF INVENTION: Sequence Determined DNA Fragments and
FTILE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 106574
LENGTH: 243
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Best Local Similarity
"herhes 5; Conserv
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NAME/KEY: peptide
; LOCATION: (1).. (240)
; OTHER INFORMATION: Ceres Seq. ID no. 13619980
US-11-056-355B-117814
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; LCCATION: (1)..(243)
; OTHER INFORMATION: Ceres Seq.
US-11-056-355B-106574
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US-11-056-355B-106574
               Sequence 117813, Application US/11056355B
publication No. US20060150283A1
GENERAL INFORMATION:
APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
APPLICANT: NIVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION UNMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
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PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 117814
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Best Local
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ORGANISM: Arabidopsis thaliana
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Arabidopsis thaliana FEATURE:
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Pred. No.
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NAME/KEY: peptide; LOCATION: (1)..(255); OTHER INFORMATION: Ceres Seq. ID US-11-056-355B-106573
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NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 117813
                                                                                                                                                                                                                                              Sequence 117812, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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LENGTH: 255
SEQ ID NO 117812
LENGTH: 255
TYPE: prt
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Best Local Similarity
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                                                                                                            APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
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CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
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TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
                                                      PRIOR APPLICATION NUMBER: 60/544,190 PRIOR FILING DATE: 2004-02-13 NUMBER OF SEQ ID NOS: 119966
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LOCATION: (1)..(243)
OTHER INFORMATION: C
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ORGANISM: Arabidopsis thaliana
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ORGANISM: Arabidopsis thaliana
FEATURE:
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Pred. No. 39;
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Pred. No.
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US-11-056-355B-68810

US-11-056-355B-68810, Application US/11056355B

; Sequence 68810, Application US/11056355B

; Publication No. US20060150283A1

; GENERAL INFORMATION:
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LOCATION: (1)..(255)
TOTHER INFORMATION: C
US-11-056-355B-117812
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TYPE: prt ORGANISM: Triticum aestivum ORGANISM: Triticum aestivum
                                                                                                         PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 68809
LENGTH: 348
                                                                                                                                                                                                     Sequence 68809, Application US/11056355B

Publication No. US20060150283A1

GENERAL INFORMATION:

APPLICANT: Brover, Vyacheslav

APPLICANT: Alexandrov, Nickolai

TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding

TITLE OF INVENTION: Polypeptides Encoded Thereby

FILE REFERENCE: 2750-1590FUS2

CURRENT APPLICATION NUMBER: US/11/056,355B

CURRENT FILING DATE: 2005-02-14
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NAME/KEY: peptide
LOCATION: (1)..(301)
OTHER INFORMATION: Ceres Seq. ID no. 12614811
NAME/KEY: peptide
LOCATION: (1)..(348)
OTHER INFORMATION: Ceres Seq.
                                                                    TYPE: prt
ORGANISM: Triticum aestivum
                                                            FEATURE:
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83.3%;
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   ID no.
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Pred. No.
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RESULT 12
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US-10-449-902-37988
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                                                                                                                                            ; ORGANISM: Arabidopsis thaliana US-10-953-349-9971
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                                                                                                                                                                                                                                                               Sequence 9971, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
CURRENT FILING DATE: 2004-09-30
                                                                                                                                                                                                            NUMBER OF SEQ ID NOS 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 9971
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 37988
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Best Local
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Best Local
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER: OF SEQ ID NOS: 56791
NUMBER: OF SEQ ID NOS: 56791
                                                                                                                                                                              TYPE: PRT
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FSRYGR 36
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83.3%;
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Pred. No. 56;
1; Mismatches
                                                                    Score 27; DB (
Pred. No. 35;
0; Mismatches
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Pred. No. 86;
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US-10-953-349-2962
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US-10-953-349-2963
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US-10-953-349-7495
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES TITLE OF INVENTION: ENCONDED THERBY FILE REFERENCE: 2750-1579PUS2 CURRENT APPLICATION NUMBER: US/10/953,349 CURRENT FILING DATE: 2004-09-30 NUMBER OF SEQ ID NOS: 40252 SOFTWARE: Patentin version 3.3 SEQ ID NO 2963 LENGTH: 209
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 7495
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GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
                                                                                                                                                                           Sequence 2962, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 2962
LENGTH: 216
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Best Local
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                                                                     APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
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TYPE: PRT
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mes 5; Conserv
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83.3%;
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                                                                                                                                                AND CORRESPONDING POLYPEPTIDES
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ORGANISM: Arabidopsis thaliana

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RESULT 16
US-10-953-349-7494
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US-10-953-349-9130
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                                               US-11-174-307B-202
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Best Local S
Matches 5
Sequence 202, Application US/11174307B Publication No. US20060143729A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 9130
LENGTH: 290
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9130, Application US/10953349 Publication No. US20060107345A1 GENERAL INFORMATION:
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: SEQUENCE-THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
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TYPE: PRT
ORGANISM: Arabidopsis thaliana
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83.3%;
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Pred. No.
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Pred. No.
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Pred. No.
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US-10-449-902-56777

Sequence 56777, Application US/10449902

Publication No. US20060123505A1

GENERAL INFORMATION:

APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution for Application for Advancement of International Science.

APPLICANT: Foundation for Advancement of International Science.

TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902

CURRENT APPLICATION NUMBER: JD 2003-05-29

PRIOR APPLICATION NUMBER: JD 2002-203269

PRIOR FILING DATE: 2002-05-30
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Best Local S
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TITLE OF INVENTION: NUCLECTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY
TITLE OF INVENTION: USEFUL FOR MODIFYING PLANT CHARACTERISTICS
FILE REFERENCE: 2750-1601PUS2
CURRENT APPLICATION NUMBER: US/11/174,307B
CURRENT FILING DATE: 2005-06-30
PRIOR APPLICATION NUMBER: 60/583,671
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR APPLICATION DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR PILING DATE: 2004-06-30
PRIOR APPLICATION DATE: 2004-06-30
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PRIOR FILING DATE: 2004-06-30
NUMBER OF SEQ ID NOS: 5544
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OTHER INFORMATION:
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OTHER INFORMATION: PFam Name: RRM_1; PFam Description: RNA recognition motif.
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NAME/KEY: misc_feature
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LOCATION:
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OTHER INFORMATION: PFam
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5; Conserv
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US-10-953-349-21500
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US-10-449-902-56777
                                         US-10-449-902-34829
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Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE THERBY
TITLE OF INVENTION: SEQUENCE THERBY
FILE REFERENCE: 2750-1579PUS2
FILE REFERENCE: 2750-1579PUS2
                                                                                                            PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 34829
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Publication No. US20060123505A1
GENERAL INFORMATION:
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SEQ ID NO 21500
LENGTH: 334
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Best Local
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     Query Match
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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SOFTWARE: PatentIn Ver. 2.1
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TYPE: PRT
                                                         TYPE: PRT
ORGANISM: Oryza sativa
                                                                                 LENGTH: 338
TYPE: PRT
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Similarity 83.3%;
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5; Conserva
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83.3%;
     87.1%; Score 27;
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Pred. No.
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RESULT 24
US-10-449-902-45028
; Sequence 45028, Application US/10449902
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PATENCHIN VERSION 3.3
SEQ ID NO 21499
LENGTH: 353
TYPE: PRT
ORGANISM: Glycine max
US-10-953-349-21499
                                                                                                                                                      ; ORGANISM: Oryza sativa US-10-449-902-33374
                                                                                                                                                                                                                                             FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: PATENTIN Vex: 2.1
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
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Publication NO. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
                                                                                                                                                                                                            SEQ ID NO 33374
LENGTH: 374
                                                                           Matches
                                                                                                               Query Match
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les 5; Conserv
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les 5; Conserv
345
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                                    1 FSRYAR 6
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83.3%;
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Pred. No.
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                                                                                             Score 27; DB 6;
Pred. No. 1e+02;
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US-11-330-403-7004
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Best Local Similarity
Thes 5; Conserve
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US-10-449-902-45028
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Best Local Similarity 83.0
Conservative
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Publication No. US20060107345A1
GENERAL INFORMATION:
NUMBER OF SEQ ID NOS: 19250
SEQ ID NO 7004
LENGTH: 478
                                                                                                                                                       Sequence 7004, Application US/11330403 Publication No. US20060159563A1 GENERAL INFORMATION:
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LENGTH: 464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE DETERMINED DNA
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: US/10/449,902
                                                       APPLICANT: Abad, Mark S.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53629)B
CURRENT APPLICATION NUMBER: US/11/330,403
CURRENT FILING DATE: 2006-01-12
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PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Glycine max
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83.3%;
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Pred. No.
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1e+02;
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US-11-353-390A-7
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SOFTWARE: PatentIn version 3.3
SEQ ID NO 7
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APPLICANT: Sticklen, Masomeh B
APPLICANT: Majbool, Shahina B
APPLICANT: Majbool, Shahina B
APPLICANT: Dale, Bruce E
TITLE OF INVENTION: TRANSGENIC PLANTS CONTAINING LIGNINASE AND CELLULASE WHICH
TITLE OF INVENTION: DEGRADE LIGNIN AND CELLULOSE TO FERMENTABLE SUGARS
FILE REFERENCE: MSU 4.1-814
                                                                                                                                                                              Sequence 7, Application US/11353390A Publication No. US20060185036A1 GENERAL INFORMATION:
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Matches
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APPLICANT: Sticklen, Masomeh B
APPLICANT: Magbool, Shahina B
APPLICANT: Magbool, Shahina B
APPLICANT: Dale, Bruce E
TITLE OF INVENTION: TRANSGENIC PLANTS CONTAINING LIGNINASE AND CELLULASE WHICH
TITLE OF INVENTION: DEGRADE LIGNIN AND CELLULOSE TO FERMENTABLE SUGARS
FILE REFERENCE: MSU 4.1-806
CURRENT APPLICATION NUMBER: US/11/353,390A
CURRENT FILING DATE: 2006-02-14
PRIOR APPLICATION NUMBER: US 60/242,408
PRIOR FILING DATE: 2000-10-20
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CURRENT FILING DATE: 2006-04-06
PRIOR APPLICATION NUMBER: US 60/242,408
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 09/981,900
PRIOR FILING DATE: 2001-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: (316)...(316)
OTHER INFORMATION: The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc feature LOCATION: (113)..(113) OTHER INFORMATION: The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Actinomyces naeslundii FEATURE:
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LOCATION: (148)..(148)
OTHER INFORMATION: The
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Pred. No.
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Pred. No. 1.3e+02;
0; Mismatches 1;
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1.3e+02;
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US-11-354-310A-7
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LENGTH: 48
TYPE: PRT
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SEQ ID NO 7
LENGTH: 488
TYPE: PRT
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Best Local
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Best Local S
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APPLICANT: Magbool, Shahina B
APPLICANT: Magbool, Shahina B
APPLICANT: Dale, Bruce E
APPLICANT: Dale, Bruce E
TITLE OF INVENTION: TRANSGENIC PLANTS CONTAINING LIGNINASE AND CELLULASE WHICH
TITLE OF INVENTION: DEGRADE LIGNIN AND CELLULOSE TO FERMENTABLE SUGARS
FILE REFERENCE: MSU 4.1-805
CURRENT APPLICATION NUMBER: US/11/354,310A
CURRENT FILING DATE: 2006-02-14
PRIOR APPLICATION NUMBER: US 60/242,408
PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 09/981,900 PRIOR FILING DATE: 2001-10-18 NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/981,900 PRIOR FILING DATE: 2001-10-18 NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.3
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                                                                                                 NAME/KEY: misc feature LOCATION: (316)...(316) OTHER INFORMATION: The
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NAME/KEY: misc_feature
LOCATION: (113)..(113)
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NAME/KEY: misc_feature
LOCATION: (113)...(113)
OTHER INFORMATION: The
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LOCATION: (148)..(148)
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                            Conservative
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83.3%;
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                            0; Mismatches
                                            Score 27; DB 7;
Pred. No. 1.3e+02;
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CURRENT APPLICATION NUMBER: US/11/174,307B
CURRENT FILING DATE: 2005-06-30
PRIOR APPLICATION NUMBER: 60/583,671
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR FILING DATE: 2004-06-30
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,651
PRIOR APPLICATION NUMBER: 60/583,651
PRIOR FILING DATE: 2004-06-30
UMMER OF SEQ ID NOS: 5544
SEQ ID NO 4272
LENGTH: 541
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                                                                                                                                                                                                 OTHER INFORMATION: GI Number: 42522036; NR Description: Gene product i OTHER INFORMATION: ubiquinone production. [Bdellovibrio bacteriovorus ; OTHER INFORMATION: >gi[39574567|emb|CAE78409.1| Gene product involved US-11-174-307B-4272
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                                                                                                                           Query Match
Best Local
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APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: UCLEDTIDE SEQUENCES AND POLYPEPTIDES ENCODED
TITLE OF INVENTION: USEFUL FOR MODIFYING PLANT CHARACTERISTICS
FILE REFERENCE: 2750-1601PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: GI Number: 24215277; NR Description: ABC1 family protein OTHER INFORMATION: kinase [Leptospira interrogans serovar Lai str. 56601] OTHER INFORMATION: >gi|24196371|gb|AAN49776.1| ABC1 family protein kinase [Leptospir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: GI Number: 45657281; NR Description: ubiquinone biosynthesis OTHER INFORMATION: protein [Leptospira interrogans serovar Copenhageni str. Fioo OTHER INFORMATION: L1-130] >gi | 45600519 | gb | AAS70004.1 | ubiquinone biosynthesis |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: GI Number: 50935257; NR Description: ABC1 family protein OTHER INFORMATION: kinase-like protein [Oryza sativa (japonica cultivar-group)] OTHER INFORMATION: >gi|23617237|dbj|BAC20904.1| ABC1 family protein kinase-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: PFam Name: ABC1; PFam Description: ABC1 family
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                                                1 FSRYAR 6
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5; Conser
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                                                                                                                         87.1%;
83.3%;
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Pred. No. 1.5e+02;
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RESULT 31 US-10-449-902-33781 ; Sequence 33781, Application US/10449902

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RESULT 33
US-11-337-300-451
; Sequence 451, Application US/11337300
; Publication No. US20060121580A1
; GENERAL INFORMATION:
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US-11-056-355B-581
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Publication No. US20060150283A
GENERAL INFORMATION:
APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nicko
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SEQ ID NO 33781
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
    APPLICANT:
                    APPLICANT: Crucell Holland B.V. APPLICANT: ter Meulen, Jan H.
                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: peptide
LOCATION: (1)..(109)
OTHER INFORMATION: Ceres Seq. ID no. 13635683
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TYPE: PRT
ORGANISM: Oryza sativa
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ORGANISM: Zea mays subsp. mays
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                                                                                                                                                                                                                                                                                                        Local Similarity
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ter Meulen, Jan H.
De Kruif, Cornelis A.
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Pred. No.
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Pred. No.
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TITLE OF INVENTION: Binding molecules against SARS-coronavirus and uses thereof; FILE REFERENCE: 0091 WO 00 ORD; CURRENT APPLICATION NUMBER: US/11/337,300; CURRENT FILING DATE: 2006-01-20; NUMBER OF SEQ ID NOS: 478; SOFTWARE: PatentIn version 3.1
SEQUENCE: 129
TYPE: DDT
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Matches
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Best Local Similarity
Matches 5; Conserv
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SEQ ID NO 31883
LENGTH: 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-30369
PRIOR FILLNG DATE: 2002-05-30
PRIOR FILLNG DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
NUMBER OF SEQ ID NOS: 56791
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Publication No. US20060123505A1
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APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILLING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
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ORGANISM: Artificial sequence
FEATURE:
FEATURE:
VARYDMATTON: Variable h
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100.0%; Pred. No.
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100.0%; Pred. No.
tive 0; Mismatc
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SEQ ID NO 286
LENGTH: 256
TYPE: PRT
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LENGTH: 219
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TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590FUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ. ID NOS: 119966
                                                 APPLICANT: Crucell Holland B.V.
APPLICANT: ter Meulen, Jan H.
APPLICANT: ter Meulen, Jan H.
APPLICANT: De Kruif, Cornelis A.
APPLICANT: Van den Brink, Edward N.
APPLICANT: Goudsmit, Jaap
TITLE OF INVENTION: Binding molecules against SARS-coronavirus and uses thereof
FILE REFERENCE: 0091 WO 00 ORD
CURRENT APPLICATION NUMBER: US/11/337,300
CURRENT FILING DATE: 2006-01-20
NUMBER OF SEQ ID NOS: 478
SOFTWARE: PatentIn version 3.1
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FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(219)
OTHER INFORMATION: Ce
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ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(219)
OTHER INFORMATION: Ceres Seq. ID no. 12566294
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100.0%; Pred. No. 1e+02;
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; TYPE: PRT; ORGANISM: Oryza sativa
US-10-449-902-31725
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US-10-449-902-30299
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                                                                                      PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 31725
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30299
LENGTH: 271
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Publication No. US20060123505A1
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT APPLICATION NUMBER: US/202-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR PILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR PILING DATE: 2002-12-11
PRIOR FILING DATE: 2002-12-11
                                                                                                                                                                                                                                                                                                                           APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
                                                                                                                                                                                                                                                          FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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100.0%; Pred. No. 1.2e+02;
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Query Match

Length 271;

Best Local Similarity

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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOCTWARE: Patentin Ver. 2.1
TYPE: DBT
LENGTH: 271
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Best Local Similarity
Matches 5; Conserva
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US-10-449-902-51604
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Publication No. US20060123505A1
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SEQ ID NO 32135
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APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205YI-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
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100.0%; Pred. No. 1.3e+02;
                                       83.9%; Score 26; DB
100.0%; Pred. No. 1.3
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; FEATURE:
, NAME/KEY: peptide
; LOCATION: (1)..(271)
; OTHER INFORMATION: Ceres Seq. ID no. 12412613
US-11-056-355B-7374
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US-11-056-355B-7374
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                                                                                                                                                        US-11-056-355B-16849
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APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
ATTILE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Sequence Determined Thereby
FILE REFERENCE: 2750-1590FUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR PILING DATE: 2004-02-13
NUMBER OF SEQ ID NO 16849
LENGTH: 271
TYPE: ATTILED TO NOS: 119966
SEQ ID NO 16849
LENGTH: 271
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SEQ ID NO 7374
LENGTH: 271
TYPE: prt
ORGANISM: Zea mays subsp. mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 7374, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 5; Conser
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                                                                            Matches
                                                                                              Query Match
Best Local Similarity
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TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
PRIOR FILING DATE: 2004-02-13
PRIOR FILING DATE: 2004-02-13
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                                                                                                                                                                      NAME/KEY: peptide
LOCATION: (1)..(271)
OTHER INFORMATION: Ceres Seq. ID no. 12566292
                                                                                                                                                                                                                                    ORGANISM: Zea mays subsp. mays FEATURE:
                                                                                                                                                                                                                                                                           TYPE: prt
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                                       1 FSRYA 5
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  FSRYA 69
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                                                                            83.9%; Score 26; DB 7;
100.0%; Pred. No. 1.3e+Cive 0; Mismatches
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Pred. No.
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                                                                                              1.3e+02;
                                                                                                                   Length 271;
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RESULT 44

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RESULT 46
US-11-330-403-19189
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, LENGTH: 334
; TYPE: PRT
; ORGANISM: Staphylococcus aureus subsp. aureus Mu50
US-11-330-403-9151
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US-11-330-403-9151
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APPLICANT: Abad, Mark S.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53629)B
CURRENT APPLICATION NUMBER: US/11/330,403
CURRENT FILING DATE: 2006-01-12
NUMBER OF SEQ ID NOS: 19250
SEQ ID NO 19189
LENGTH: 334
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION: APPLICANT: Abad, Ma
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                                                                                                                                                                          Sequence 19189, Application US/11330403
Publication No. US20060159563A1
GENERAL INFORMATION:
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Publication No. US20060159563A1
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Matches 5; Conserv
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TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Genes and Uses for Plant Improvement FILE REFERENCE: 38-21 (33629)8 CURRENT APPLICATION NUMBER: US/11/330,403 CURRENT FILING DATE: 2006-01-12 NUMBER OF SEQ ID NOS: 19250
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ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(311)
OTHER INFORMATION: Ceres Seq. ID no. 12412611
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                                                                                                                                                                                                                                                                                                                                                                                     83.9%;
Local Similarity 83.3%;
es 5; Conservative
                                                                                                                                                                                                                                                                                                                     103
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100.0%; Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Query Match
Best Local Similarity
"hes 5; Conservi
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                                                   ; TYPE: PRT
; ORGANISM: Homo
US-11-359-554-3
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                                                                                                                        SOFTWARE: FastSEQ
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/11359554
Publication No. US20060154293A1
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Best Local Similarity
Query Match
Best Local Similarity
                                                                                                                                                          PRIOR APPLICATION NUMBER: US/10/311,591
PRIOR FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: US 60/213,474
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/277,612
PRIOR PILING DATE: 2001-03-22
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                      FILE REFERENCE: 004974.00929
CURRENT APPLICATION NUMBER: US/11/359,554
CURRENT FILING DATE: 2006-02-23
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Xiao, Yonghong
TITLE OF INVENTION: Regulation of Human Prostasin-Like
TITLE OF INVENTION: Serine Protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/280,509
PRIOR FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: US 60/311,055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/11/404,745
CURRENT FILING DATE: 2006-04-14
PRIOR APPLICATION NUMBER: US/10/109,616
PRIOR FILING DATE: 2002-03-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Allen, Keith D.
TITLE OF INVENTION: TRANSGENIC MICE CONTAINING CHANNEL
TITLE OF INVENTION: ACTIVATING PROTEASE 1 (CAP1) GENE DISRUPTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: R-490
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                                                                                                       LENGTH: 343
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 83.9%;
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 Mismatches

Score 26;
Pred. No.
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Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
 1.6e+02;
                   DB 7;
                 Length 343;
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APPLICANT: Handfield, Martin
APPLICANT: Handfield, Martin
APPLICANT: Handfield, Meffrey
APPLICANT: Hillman, Jeffrey
APPLICANT: Progulske-Fox, Ann
TITLE OF INVENTION: Identification of Actinobacillus actinomycetemcomitans
TITLE OF INVENTION: Antigens for Use in the Diagnosis, Treatment, and Monitoring
TITLE OF INVENTION: Antigens for Use in the Diagnosis, Treatment, and Monitoring
TITLE OF INVENTION: Antigens for Use in the Diagnosis, Treatment, and Monitoring
TITLE OF INVENTION: Antigens for Use in the Diagnosis, Treatment, and Monitoring
TITLE OF INVENTION: Antigens for Use in the Diagnosis, Treatment, and Monitoring
TITLE OF INVENTION: SALID INVENTION INVE
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US-10-471-571A-4440
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US-11-333-747A-22
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, NAME/KEY: MISC FEATURE
; LOCATION: (1). (348)
; OTHER INFORMATION: conserved hypothetical
US-10-471-571A-4440
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CURRENT APPLICATION NUMBER: US/10/471,571A
CURRENT FILING DATE: 2003-09-12
PRIOR APPLICATION NUMBER: GB-0107661.1
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 5642
SOFTWARE: SeqWin99, version 1.03
SEQ ID NO 4440
LENGTH: 348
TYPE: PRT
ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22, Application US/11333747A Publication No. US20060160159A1 GENERAL INFORMATION:
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                                                                                                                         Query Match
Best Local Similarity
Matches 5; Conserv
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ative 0; Mismatches 0;
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Search completed: August 29, 2006, 11:29:12 Job time: 4.75248 secs

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ABB95971925

AAG18086

ADX91925091

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                                                                                                  The present invention relates to motoneuronotrophic factor peptide analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are useful for promoting motor neuron viability and axon degeneration, target muscle reinnervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New motoneuronotrophic factor peptide analogue enhancing the viability motor neurons, useful for target muscle reinnervation, treating peripheral nerve injuries or neurodegenerative diseases and in wound
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                                                                                                                                                                                                                                                                                                                                                          Claim
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                                                                         represents a MNTF motoneuronotrophic factor peptide analogue
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                                                                                                                                                                                                                                                                                                                                                       NO 6; 40pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Sequence 13 AA;

RESULT 4 AAO29914 ID AAO2

AAO29914 standard; protein; 33 AA.

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12

WMLSAFS

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RESULT 3
AAWS9046
ID AAWS
XX AAWS
AC AAWS
XX Huma
XX Huma
XX Moto
KW Moto
CO 2-4
PR 12-4
PR 12
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Best Local S
Matches 7
                                                        Query Match
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Matches 7
                                                                                                                                                                                                                          This sequence, represents a fragment of a novel human motoneurotrophic factor, MNTF1-F6. Such factors are used to promote regeneration of the axon of a motoneurone, to diagnose and treat motoneurone disease in a mammal or to accelerate wound healing whilst concomitantly minimising or inhibiting scar tissue and/or keloid formation in an area associated with a wound. For promoting axonal regeneration, the polypeptide is administered at a concentration of 5 ng-50 mg, whereas for inhibiting hereditary motoneurone disease, the dosage is 5-100 (especially 30-50) ng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-SEP-1996;
15-NOV-1996;
12-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Motoneurotrophic factor MNTF1-F3 and MNTF1-F6 - useful for motoneuron regeneration, diagnosing or treating motoneuron disease and to accele, wound healing without scar formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-1998
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                                                                                                                                                        Sequence 33
                                                                                                                                                                                                          per kg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Fig 2B; 78pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chau RMW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human MNTF1-F6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1998-230703/20
                                                                                                                                                                                                             body weight
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                                                     Similarity
7; Conserv
        1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BIOTECH INC
                                                                                                                                                          AA;
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                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96US-0026792P.
96US-00751225.
97US-00928862.
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                                                                                100.0%;
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                                                                                Score 38;
Pred. No.
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                                                          Mismatches
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                                                                                                        Length
                                                          Indels
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RESULT 5
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AC ADQ9
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DT 21-C
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MNTF
XX
MOTO
KW MOTO
KW Neux
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                                                                                                                                                                                                                                                                                                                                                                                           peripheral nerve injuries, musculoskeletal disorders, spinal cord injuries, head injuries, strokes, neuromuscular degenerative diseases, amyotrophic lateral sclerosis, spinal muscular atrophy, peripheral neuropathy, diabetic peripheral neuropathy, peripheral neuropathy resulting from AIDS or radiation therapy for cancer, multiple sclerosis, muscular dystrophy, inhibition of scar formation, myasthenia gravis and sensory neuronal function disorders. The present sequence is human MNTF1-F6 protein used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method for promoting the survival, growth proliferation or maintenance of mammalian neurons by administering motoneuronotrophic factors (MNTF). The method is useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Promoting the survival, growth, proliferation or maintenance of mammalian neurons by administering motoneuronotrophic factors, useful for treating musculoskeletal and neurodegenerative disorders and spinal cord injuries.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   head injury; stroke; neuromuscular degenerative disease; nootropic; amyotrophic lateral sclerosis; peripheral neuropathy; neuroprotective; muscular dystrophy; AIDS; spinal muscular atrophy; multiple sclerosis;
 Neuropeptide-Agonist;
               motoneuronotrophic; motor neuron; Nootropic; Neuroprotective; Vulnerary;
                                             MNTF1-F6 33 mer motoneuronotrophic factor peptide analogue.
                                                                                 21-OCT-2004
                                                                                                                                                ADQ90195 standard; peptide; 33
                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Fig 2B; 90pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peripheral nerve injury; musculoskeletal disorder; spinal cord inju
head injury; stroke; neuromuscular degenerative disease; nootropic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; motoneuronotrophic factor; MNTF; therapy; radiation therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        motoneuronotrophic factor
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                                                                                                                                                                                                                                  WMLSAFS 18
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ω
                                                                                                                                                                                                                                                                                                                                                               A,
                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gravis; sensory neuronal
                                                                                                                                                                                                                                                                                                              100.0%;
 axon
degeneration;
                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                              Score 38; DB
Pred. No. 2.8;
                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        atrophy; multiple sclerosis;
function disorder; vulnerary;
                                                                                                                                                                                                                                                                                                                            6
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                                                                                                                                                                                                                                                                                                                            Length 33;
                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          growth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      injury;
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to motoneuronotrophic factor peptide analogue, where the peptide analogue enhances the viability of motor neurons. The methods and compositions of the present invention are useful for promoting motor neuron viability and axon degeneration, target muscle reinnervation, treating peripheral nerve injuries, treating neurodegenerative disease and in wound healing. The present sequence represents MNTF1-F6 33 mer motoneuronotrophic factor peptide analogue.
                                                    Venter JC,
                                                                                                                                                              23-MAR-2000;
11-JUL-2000;
                                                                                                                                                                                                                                                                                                   27-SEP-2001
                                                                                                                                                                                                                                                                                                                                                     WO200171042-A2
                                                                                                                                                                                                                                                                                                                                                                                                       Drosophila melanogaster.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Drosophila melanogaster polypeptide SEQ ID NO 16893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New motoneuronotrophic factor peptide analogue enhancing the viability motor neurons, useful for target muscle reinnervation, treating peripheral nerve injuries or neurodegenerative diseases and in wound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JAN-2003; 2003US-0441772P
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                                                                                                                                                                                                                                            23-MAR-2001; 2001WO-US009231.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila; developmental biology; cell signalling; insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABB63367 standard; protein; 1760
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Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL040-ABL16175) and the encoded proteins (ABE9737-ABB72072). The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly
The invention describes a method of identifying compounds that inhibit the activity of, or that interact with a protein essential for Drosophila larval viability comprising expressing in a recombinant host a DNA molecule to produce a protein essential for larval viability. The method is useful for identifying compounds with insecticidal activiti.
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                                                                                                                      Identifying inhibitors of activity of proteins essential for Drosophila larval viability comprises expressing in a host a protein essential for larval activity and identifying compounds that inhibit or interact with
                                                                                                                                                                                                                    Stam L,
                                                                                                                                                                                                                                                                                                                                                                                    Drosophila melanogaster
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                                                                                Claim 1; Page 120-126; 169pp; English
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sed rape; soybean; vegetable crop; fruit.
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barley; millet; turf; cotton; sugarcane; suga
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Niknejad A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         identified are useful as insecticides in crops such as maize, wheat, oats, rye, sorghum, rice, barley, millet, turf, cotton, sugarcane, sughet, oilseed rape, soybeans, vegetable crops and fruits. This is the amino acid sequence of a fruit fly larval viability associated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1760 AA;
                                                                                                                                       Screening and/or diagnosing cardiovascular disorder in subject involves detecting and/or quantifying level of polypeptide in biological sample from subject and comparing with control sample.
                                                                                                                                                                                                                                                                                     08-AUG-2003; 2003US-0493599P
08-AUG-2003; 2003US-0493836P
08-AUG-2003; 2003US-0493867P
08-AUG-2003; 2003US-0493985P
                                                                                                                                                                                                                                                                                                                                                                                                                                              dermatological disease; musculoskeletal disease; Cardiant; Vasotropic; Hepatotropic; Litholytic; Immunosuppressive; Gastrointestinal-Gen.; Antiinflammatory; Antidiabetic; Dermatological; Antiemetic; Laxative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              emesis; constipation; diarrhea; cardiovascular disease; immune dendocrine disease; gastrointestinal disease; metabolic disorder;
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(NOVS ) NOVARTIS A
(NOVS ) NOVARTIS PI
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85.7%;
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se K, Saudrais
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The invention relates to a method of screening and/or diagnosing a cardiovascular disorder (CD) in a subject which comprises detecting and/or quantifying the level of a polypeptide in a biological sample from the subject and comparing the level to that of control sample. The method is useful for screening, diagnosing and treating coronary artery disease, biliary cirrhosis, gallstones, celiac disease, irritable bowel syndrome, diabetes, scleroderma, nausea, vomiting, constipation and diarrhea. The method is rapid and efficient. The present sequence represents a cardiovascular disorder plasma protein tryptic fragment.

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NO 236;

349pp; English.

Sequence 17 AA;

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                                                                                      CC lubricating molety and has a polypeptide sequence comprising 1-76 repeats CC of a motif having at least 50% identity to the sequence KEPAPTT CC (AAB29774). The invention also relates to a nucleic acid encoding a human CC MSF-derived tribonectin; a biocompatible composition comprising a human CC tribonectin for inhibiting tissue adhesion formation; and a method of CC diagnosing osteoarthritis or a predisposition to osteoarthritis by CC measuring the amount of MSF or its fragment in a biological sample of a CC indicates the presence of or predisposition to developing osteoarthritis. CC indicates the presence of or predisposition to developing osteoarthritis. CC indicates the presence of or predisposition to developing osteoarthritis. CC osteoarthritis, where they may be used for lubricating mammalian joints, CC such as articulating joints of humans, dogs or horses. The tribonectin, CC when formulated as a membrane, foam, gel or fibre, is useful for CC inhibiting adhesion between two surfaces such as the injured tissues of a CC mammal, where the injury is caused by a surgical insertion or trauma, or can artificial device e.g., an orthopaedic implant. In particular, one of the surfaces is pericardial tissue. DNA encoding a tribonectin may be used in gene therapy. The present sequence represents a fragment of a human MSF-derived tribonectin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a human tribonectin which is a product of alternative splicing of the human MSF (megakaryocyte stimulating factor)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel tribonectin polypeptide useful as lubricant for treating osteoarthritis, comprises O-linked lubricating moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 1; 47pp; English
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Query Match

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RESULT 10
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                                                                                                                                                      ABB43398 to ABB44239. The proteins can have activities based on the citissues and cells the genes are expressed in. Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory; antityroid; antiarlergic; antibacterial; antiviral; cd antiinflammatory; antityroid; antiallergic; antibacterial; antiviral; concurrence; antiprotective; cardiant; thrombolytic; coagulant; compositic; coagulant; compositic; coagulant; compositic; coagulant; compositic; coagulant; composities and antiangiogenic. The motivating medical conditions and diagnosing pathological conditions. Cc polynucleotides and polypeptides can be used for preventing, treating or cmeliorating medical conditions and diagnosing pathological conditions. Cc polynucleotides, polypeptides, antibodies, agonists and antagonists from cc the present invention may be used to treat immune disorders by activating cor inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autodimmune cells, altergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnosis; cytostatic; proliferative; vulnerary; immunomodulator;
antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                    inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. ARC78449 to ARC78457 and AAB44240 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 11; Page 2133; 2352pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing \mathbf{e}.\mathbf{g}. cancer.
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N-PSDB; AAC78216.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC
                            present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer associated protein sequence SEQ ID NO:1452
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                                                                                                                                             The present invention provides the protein and coding sequences of human superficial zone protein (SZP). The protein is involved in the lubrication of joints, and the sequences can be used in the treatment of degenerative joint conditions or to delay symptoms of a degenerative joint condition, e.g. osteoarthritis, rheumatoid arthritis, gout, psoriatic arthritis, reactive arthritis, viral or post viral arthritis, spondylarthritis, juvenile arthritis, synovitis, tendonitis, systemic lupus erythematosus, CACP, osteoporosis or trauma. The present sequence is the protein encoded by the human 3' cartilage SZP cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New purified superficial zone protein (SZP) polypeptides, useful treating degenerative joint conditions, e.g. osteoarthritis, rhewarthritis, gout, spondylarthritis, synovitis, tendonitis, lupus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SZP; superficial zone protein; cartilage; lubrication; human; degenerative joint condition; arthritis; osteoporosis; trauma; CACP; chondroitin sulphate substitution consensus; antiarthritic; antirheumatic; osteopathic; antigout; antiinflammatory; dermatological;
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N-PSDB; AAL49079.
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Dixon EP;
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Local Similarity nes 6; Conserv
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811 WMLSPFS

1 WMLSAFS 7

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RESULT 12
AAB29778
JID AAB29
XX AAB29
XX AAB29
XX LIVE ALL AAB29
XX AAB29
XX LIVE ALL AAB29
XX AAB29
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XX Human
XX Human
XX Homo
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XX AC
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                                                                                                                                                                                    cc diagnosing osteoarthritis or a predisposition to osteoarthritis by companied to the amount of MSF or its fragment in a biological sample of a mammal, wherein an increased amount of MSF compared to a control comment in the presence of or predisposition to developing osteoarthritis. Cc The tribonectin and DNA encoding it are useful in the treatment of costeoarthritis, where they may be used for lubricating mammalian joints, cush as articulating joints of humans, dogs or horses. The tribonectin, cc when formulated as a membrane, foam, get or fibre, is useful for combibiting adhesion between two surfaces such as the injured tissues of a mammal, where the injury is caused by a surgical insertion or trauma, or can artificial device e.g., an orthopsedic implant. In particular, one of the surfaces is pericardial tissue. DNA encoding a tribonectin may be used in gene therapy. The present sequence represents a substantial coportion of a human MSF-derived tribonectin
                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tubricating molety and has a polypeptide sequence comprising 1-76 repeats of a motif having at least 50% identity to the sequence KEPAPTI (ABE29774). The invention also relates to a nucleic acid encoding a human MSF-derived tribonectin; a biocompatible composition comprising a human tribonectin for inhibiting tissue adhesion formation; and a method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human tribonectin; MSF; megakaryocyte stimulating factor; alternative splicing; joint boundary lubricant; O-linked oligosaccharide; osteoarthritis; tribosupplementation; tissue adhesion inhibition; friction coefficient reduction; gene therapy; antiarthritic; osteopathic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human MSF-derived tribonectin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB29778;
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                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a human tribonectin which is a product of alternative splicing of the human MSF (megakaryocyte stimulating factor)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 1; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel tribonectin polypeptide useful as lubricant for treating osteoarthritis, comprises O-linked lubricating moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-024673/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jay GD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RHOD-) RHODE ISLAND HOSPITAL LIFESPAN PARTNER.
                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The tribonectin has at least one O-linked oligosaccharide
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l Similarity
6; Conserv
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      Conservative
                                                                                                                                       B
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                                      86.8%;
      0;
                                      Score 33;
Pred. No.
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             Mismatches
                                         9.3e+02;
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                                                                     Length 902
             Indels
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RESULT 13
ADQ67713
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                                                                                                                                                                                                             Matches
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Best Local (
                                                                                                                                                                                                                                                                                              The invention relates to 2495 novel polynucleotides (I) and their encoded polypeptides, sequences hybridizing to these nucleotides, sequences encoding partial polypeptides and sequences having 70% or 90% identity to the nucleotide and protein sequences. The nucleotides and polypeptides are useful as diagnostic markers or therapeutic target for the diseases or morbid states. They are also useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers. This sequence corresponds to a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteopathic; neuroprotective; nootropic; antiparkinsonian; cytostatic; gene therapy; diagnostic marker; morbid state; osteoporosis; neurological disease; Alzheimer's disease; Parkinson's disease; dementia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-JAN-2003; 2003JP-00102206.
09-MAY-2003; 2003JP-00131392.
   lubricin;
                           PRG4-Lub:1
                                                  02-JUN-2005
                                                                          ADZ03664;
                                                                                                ADZ03664 standard; protein; 981 AA
                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 4874; 2449pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel 2495 cDNA, useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-OCT-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REAS-)
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                                                                                                                                                               700 WMLSPFS 706
                                                                                                                                                                                     1 WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RES ASSOC BIOTECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ67406.
                                                                                                                                                                                                                                                              933 AA;
                                                                                                                                                                                                                                                                                       of the invention.
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antiarthritic; osteopathic; antirheumatic; cytostatic;
                         protein.
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                                                                                                                                                                                                               Conservative
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                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isono Y,
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                                                                                                                                                                                                                          86.8%;
85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  933
                                                                                                                                                                                                             0,
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Pred. No. 9.7e+02;
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                                                                                                                                                                                                             Mismatches
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ADZ03668
ID ADZ03
XX ADZ03
XX ADZ03
XX DZ03
XX DZ03
XX DZ07
XX DZ07
XX DZ07
XX PRG4-
XX Lubri
KW cardi
KW cardi
KW neopl)
XX Synth
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PD 24-FE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated lubricin protein. A lubricin containing composition is useful for treating a subject which involves administering same to a tissue (e.g. cartilage, synovium, meniscus, tendon, peritoneum, pericardium, or pleura (preferably cartilage) of the subject (e.g. mouse, rat, cat, dog, horse or human (preferably human)). The above method further involves providing an anasthetic, antinflammatory drug and antibiotic to the subject, aspirating fluid from the subject, washing tissue of the subject and imaging tissue of the subject. The protein is useful for treating osteoarthritis or rheumatoid subject. The protein is useful for treating osteoarthritis or rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, peritoneum, peritoardium or pleura. An anti-lubricin antibody is useful for treating cancer e.g. synoviomas. The present sequence represents the amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel recombinant lubricin useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, peritoneum, pericardium or pleura.
                                                                                                                lubricin; antiarthritic; osteopathic; antirheumatic; cytostatic; cardiant; respiratory-gen.; osteoarthritis; rheumatoid arthritis neoplasm; musculoskeletal disease; immune disorder; inflammation
                                                                                                                                                                                                                      02-JUN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 981 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer e.g. synoviomas. The present sequence of the PRG4-Lub:1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 11; SEQ ID NO 7; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Flannery CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-AUG-2003; 2003US-0495741P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cardiant; respiratory-gen.; osteoarthritis; rheumatoid arthritis; neoplasm; musculoskeletal disease; immune disorder; inflammation.
                                                 WO2005016130-A2
                                                                                                                                                                                   PRG4-Lub:2 protein
                                                                                                                                                                                                                                                                                        ADZ03668 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                          1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                          754
                                                                                                                                                                                                                                                                                    protein; 1007
                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             86.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1e+03;
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RESULT 16
ADZ03672
ID ADZ03
XX
AC ADZ03
XX
DT 02-JU
XX
Cardi
KW lubri
KW cardi
KW neopl
XX
NOS Syntl
XX
NOS Syntl
XX
NOS 13-AI
XX
PF 13-AI
XX
PR (AMH-
XX
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XX
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       administering same to a tissue (e.g. cartilage, synovium, meniscus, tendon, peritoneum, pericardium, or pleura (preferably cartilage)) of the subject (e.g. mouse, rat, cat, dog, horse or human (preferably human)). The above method further involves providing an anasthetic, antiinflammatory drug and antibiotic to the subject, aspirating fluid from the subject, washing tissue of the subject and imaging tissue of the subject. The protein is useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articular pericardium or pleura. An anti-lubrician antibody is useful for treating cancer e.g. synoviomas. The present sequence represents the amino acid sequence of the PRG4-Lub:2 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel recombinant lubricin useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, peritoneum, pericardium or pleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1007 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated lubricin protein. A lubricin containing composition is useful for treating a subject which involves
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 11; SEQ ID NO 11; 19pp; English
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                                                                                                                                                                                                                     Synthetic
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                                                                                                    14-AUG-2003; 2003US-0495741P
                                                                                                                                 13-AUG-2004; 2004WO-US026508
                                                                                                                                                                                           WO2005016130-A2
                                                                                                                                                                                                                                                    neoplasm;
                                                                                                                                                                                                                                                                  lubricin; antiarthritic; osteopathic; antirheumatic; cytostatic;
cardiant; respiratory-gen.; osteoarthritis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                             PRG4-Lub:3 protein
                                                                                                                                                                                                                                                                                                                                                                                                    ADZ03672 standard;
               2005-272808/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            774 WMLSPFS 780
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                                            CR,
                                                                                                                                                                                                                                                    musculoskeletal disease; immune disorder; inflammation
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                                            Corcoran CJ,
                                                                                                                                                                                                                                                                                                                                                                                                  protein;
                                                                                                                                                                                                                                                                                                                                          entry)
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Pred. No.
                                            Freeman
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                                            BA,
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                                            Racie
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated lubricin protein. A lubricin containing composition is useful for treating a subject which involves administering same to a tissue (e.g. cartilage, synovium, meniscus, tendon, peritoneum, pericardium, or pleura (preferably cartilage)) of the subject (e.g. mouse, rat, cat, dog, horse or human (prefarably human)). The above method further involves providing an anasthetic, antiinflammatory drug and antibiotic to the subject, aspirating fluid from the subject, washing tissue of the subject and imaging tissue of the subject. The protein is useful for treating osteoarthritis or relumnatoid arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, periconeum, pericardium or pleura. An anti-lubricin antibody is useful for treating cancer e.g. synoviomas. The present sequence represents the amino acid sequence of the PRG4-Lub:3 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel recombinant lubricin useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, peritoneum,
                                                           rheumatoid arthritis, or as lubricants, articular supplements for synovial joint pericardium or pleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1038 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; SEQ ID NO 15; 19pp; English
                                Claim 11; SEQ ID NO 19; 19pp; English
                                                                                                                                      WPI; 2005-272808/28.
N-PSDB; ADZ03675.
                                                                                                                                                                                   Flannery CR,
                                                                                                                                                                                                                                              14-AUG-2003; 2003US-0495741P
                                                                                                                                                                                                                                                                            13-AUG-2004; 2004WO-US026508
                                                                                                                                                                                                                                                                                                           24-FEB-2005
                                                                                                                                                                                                                                                                                                                                        WO2005016130-A2
                                                                                                                                                                                                                                                                                                                                                                                                   neoplasm;
                                                                                                                                                                                                                                                                                                                                                                                                                    cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                lubricin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRG4-Lub:4 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-JUN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADZ03676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADZ03676 standard; protein; 1069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pericardium or pleura.
                                                                                                           Novel recombinant lubricin useful for treating osteoarthritis or
                                                                                                                                                                                                                 (AMHP ) WYETH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      805 WMLSPFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                   antiarthritic; osteopathic; antirheumatic; cytostatic;
respiratory-gen.; osteoarthritis; rheumatoid arthritis
musculoskeletal disease; immune disorder; inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                    Corcoran
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                    Freeman BA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                            joints,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1e+03;
                                                                                            anti-adhesive agent or intra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 9;
                                                                            meniscus, tendon, peritoneum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 1038;
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invention relates to an isolated lubricin protein. A lubricin

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Matches
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The invention relates to an isolated lubricin protein. A lubricin containing composition is useful for treating a subject which involves administering same to a tissue (e.g. cartilage, synovium, meniscus, tendon, peritoneum, pericardium, or pleura (preferably cartilage)) of the subject (e.g. mouse, rat, cat, dog, horse or human (preferably human)). The above method further involves providing an anasthetic, antiinflammatory drug and antibiotic to the subject, aspirating fluid from the subject, washing tissue of the subject and imaging tissue of the subject. The protein is useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        subject (e.g. mouse, rat, cat, dog, horse or human (preferably human)). The above method further involves providing an anasthetic, antiinflammatory drug and antibiotic to the subject, aspirating fluid from the subject, washing tissue of the subject and imaging tissue of the subject. The protein is useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or intra articular supplements for synovial joints, meniscus, tendon, periconeum, pericardium or pleura. An anti-lubricin antibody is useful for treating cancer e.g. synoviomas. The present sequence represents the amino acid sequence of the PRG4-Lub:4 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                        Novel recombinant lubricin useful for treating osteoarthritis or rheumatoid arthritis, or as lubricants, anti-adhesive agent or in articular supplements for synovial joints, meniscus, tendon, per
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lubricin; antiarthritic; osteopathic; antirheumatic; cytostatic;
cardiant; respiratory-gen.; osteoarthritis; rheumatoid arthritis;
neoplasm; musculoskeletal disease; immune disorder; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   subject (e.g. mouse
                                                                                                                                                                                                                                                                                                                                           Claim 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                pericardium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADZ03679.
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                                                                                                                                                                                                                                                                                                                                        SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein.
                                                                                                                                                                                                                                                                                                                                                                                                or pleura.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 1100 AA.
                                                                                                                                                                                                                                                                                                                                     23; 19pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Freeman BA,
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Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                        e agent or intra
tendon, peritoneum,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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The present sequence is that of novel human extracellular messenger (EXMES) Incyte ID NO: 7513017CD1 polypeptide. The protein shows homology to human megakaryocyte stimulating factor. The invention provides EXMES polynucleotides and polypeptides, as well as expression vectors, host cells, antibodies, agonists and antagonists, and methods for diagnosing, treating or preventing disorders associated with aberrant expression of

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RESULT 19
ADX67911
ID ADX677
XX
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ADX67917
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ADX67917
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                                                                                                                                                                                                                                      New extracellular messengers and nucleic acids, useful for diagnosing, treating or preventing e.g. allergies, anemia, Crohn's disease, diabet mellitus, myasthenia gravis, osteoporosis, psoriasis, cancers or
                                                                                                                                                                                                                                                                                                                                                                                                                                       Elliott VS
Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-AUG-2002; 2002US-0400810P.
19-SEP-2002; 2002US-0412197P.
04-OCT-2002; 2002US-0416004P.
08-NOV-2002; 2002US-0424862P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   muscular-gen.; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antiasthmatic;
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                                                                                                                                                                                                                  autoimmune thyroiditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-2003; 2003WO-US024084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004013292-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  extracellular messenger (EXMES) polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              vs,
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                                                                                                                                                                                                                                                                                                                                             ADK67916.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             atic; antiinflammatory; antidiabetic; neuroprotective;
gen.; antiarthritic; osteopathic; hepatotropic; antipsoriatic;
fungicide; antiparasitic; protozoacide; antihelminthic;
                                                                                                                                                                                                                                                                                                                                                                                                                                          TW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene
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1, Emerling BM,
                                                                                                                                                                       IJ
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according to identification method"
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note= "Spans residues 1 to 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy
                                                                                                                                                                     NO 4; 165pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                       UK, Swarnakar A,
, Lindquist EA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 33; DB Pred. No. 1.2e 0; Mismatches
                                                                                                                                                                       English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EXMES;
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.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Chawla NK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 1100;
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                                                                                                                                                                                                                                                                                                                                                                                                                      s JP;
K. Ramkumar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for treating
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RESULT 20
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Best Local S
Matches 6
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17-JUL-2000;
03-AUG-2000;
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                           proteins from a variety of organisms, including human, dog, cat, ho cow, pig, hamster, monkey, macague, yeast, bacteria, fruit fly, sea urchin and tomato. These were derived from expressed sequence tags from the organism of interest. They can be used in diagnostics, forensics, gene mapping, identification of mutations, to assess biodiversity and for nutritional purposes. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; sheep; pig; cow; fruit fly; yeast; hamster; macaque; horse; tomato; monkey; dog; sea urchin; expressed sequence tag; EST; diagnostics; forensic test; gene mapping; genetic disorder; biodiversity;
                                                                                                                                                                                                                                                                                                                      Isolated polypeptide for treatment of antibodies and research use.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200154477-A2
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                                                                                                                                                                                                        The present invention provides the protein and coding sequences of novel
                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-476164/51.
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                                                                                                                                                                                                                                                               Page 1198-1201; 1275pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 2000US-00491404.
; 2000US-00617746.
; 2000US-00631451.
; 2000US-00663870.
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J, W
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                              diagnostics, raising
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Best Local S
Matches 6
                                The present sequence is that of novel human extracellular messenger (EXMES) Incyte ID NO: 7513018CD1 polypeptide. The protein shows homology to human megakaryocyte stimulating factor. The invention provides EXMES polymucleotides and polypeptides, as well as expression vectors, host cells, antibodies, agonists and antagonists, and methods for diagnosing, treating or preventing disorders associated with aberrant expression of EXMES, especially autoimmune and inflammatory disorders, cell
                                                                                                                                                   New extracellular messengers and nucleic acids, useful for diagnosing, treating or preventing e.g. allergies, anemia, Crohn's disease, diabet mellitus, myasthenia gravis, osteoporosis, psoriasis, cancers or autoimmune thyroiditis.
                                                                                                                                                                                                                                                                                                                        02-AUG-2002; 2002US-0400810P.
19-SEP-2002; 2002US-041212P9.
04-OCT-2002; 2002US-0416004P.
08-NOV-2002; 2002US-0424862P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; extracellular messenger; EXMES; respiratory-gen.; antiallergic; antiasthmatic; antiinflammatory; antidiabetic; neuroprotective; muscular-gen.; antiarthritic; osteopathic; hepatotropic; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1299
                                                                                                                               Claim
                                                                                                                                                                                                                                                             Richardson
                                                                                                                                                                                                                                                                         Elliott VS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytostatic; gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK67912;
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                                                                                                                                                                                                               2004-157116/15.
DB; ADK67917.
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6; Conserv
                                                                                                                                                                                                                                                                                                  INCYTE CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fungicide; antiparasitic; protozoacide;
                                                                                                                              SEQ ID NO 5; 165pp; English.
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W, Emerling
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/note= "Spans residues 1 to 18, 20,
according to identification method"
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ng BM,
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Pred.
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Chawla NK
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proliferative disorders and endocrine disorders, e.g. adult respiratory distress syndrome, allergy, anaemia, erythroblastosis foetalis, Crohn's disease, diabetes mellitus, myasthenia gravis, osteoarthritis,

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RESULT
ADK6581
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                                         The invention relates to a method of determining the angiogenic index of a tissue or cell sample comprising assessing, in a sample, the expression levels of one or more differentially-expressed gene from any of 34 DNA sequences, given in the specification, where the levels are indicative of the angiogenic index. The methods and compositions of the present invention are useful for diagnosing, preventing and/or treating cancer, coronary artery disease, myocardial ischemia or coronary artery disease, myocardial ischemia or coronary arteriosclerosis. They can also be used in research, drug discovery and forensic medicine involving angiogenesis. This sequence corresponds to
                                                                                                                                                                                                                                                                                                                                                                                               Determining the angiogenic index of a tissue or cell sample using expression levels of differentially expressed genes, useful for diagnosing or treating cancer, coronary artery disease, myocardial ischemia and/or arteriosclerosis.
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10-JUN-2002; 2002US-00164595.
16-AUG-2002; 2002US-0403649P.
03-JAN-2003; 2003US-0437746P.
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                                                                                                                                                                                                                                                                                                                                             Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-FEB-2003; 2003WO-US003848.
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19
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                    the differentially expressed proteins of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORIGENE TECHNOLOGIES INC
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                                                                                                                                                                                                                                                                                                                                          SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cardiant; vasotropic; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                       58; 296pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fan W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jay G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein ANH0316.
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                          invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1320 AA;
            New human mega-karyocyte stimulating factors - for treating immune deficiencies, cancer, exposure to radiation or drugs, bacterial and
                                                                                                                                                                    Region
                                                                                                                                                                                Region
                                                                                                                                                                                             Region
                                                                                                                                                                                                         Region
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                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                     stability;
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02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                       AAR26049
                                N-PSDB;
                                     WPI; 1992-284660/34
                                                                           18-JAN-1991;
10-SEP-1991;
                                                                                               17-JAN-1992;
                                                                                                            06-AUG-1992
                                                                                                                                                       Region
                                                                                                                                                                                                                                  Region
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                                                                                                                                                                                                                                                                                                                           Megakaryocyte
                                                                                                                                                                                                                                                                                                                                        MSF precursor
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            deficiencies, cancer,
                                                                                                                        WO9213075-A1
                                                               (GEMY ) GENETICS INST INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 6; Conserv
                                AAQ27223.
                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                   Clark
                                                                                                                                                                                                                                                                                                                     /te colony stimulating
proteolytic cleavage;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                     (revised)
(first en
                                                                            91US-00643502
91US-00757022
                                                                                               92WO-US000433
                                                                                                                              /label= Fv
                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                  SC,
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                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   86.8%;
                                                                                                                                                      el= Exon_XI
.1404
                                                                                                                                                                                l= Exon_IX
.1331
                                                                                                                                                                   l= Exon_X
.1373
                                                                                                                                                                                                         l= Exon_VI
.1212
                                                                                                                                                                                                                                                 . 157
                                                                                                                                                                                                                       .1141
                                                                                                                                                                                                                                                                 l=_Exon_II
                                                   Jacobs
                                                                                                                                                                                             .1266
                                                                                                                                                                                                                                                                              = Exon_I
                                                                                                                                                                                                                                                     Exon_III
                                                                                                                                               Exon_XII
                                                                                                                                                                                                 Exon_VIII
                                                                                                                                                                                                                           Exon_V
                                                                                                                                   Exon_VII
                                                                                                                                                                                                                                         Exon_IV
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                     factor; secretion signal;
adhesion; alternative spl
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                                                   Hewick RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 7;
                                                   Gesner
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 1320;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                     splicing
                                                                                                                                                                                                                                                                                                                            meg-CSF;
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              viral
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infections, etc.

Claim 1, 2 and 3; Fig 1; 87pp; English

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ARBSULT 24
AAB29773
ID AAB29
XX AAB29
XX AAB29
XX AB29
XX AB29
XX AB29
XX AB29
XX AB29
XX Human
XX Human
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XX Homo
XX AC20
XX A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence given is a full length translation from the megakaryocyte cot stimulating factor (MSP) precursor. The sequence covered by exons II. III can and IV encodes megakaryocyte stimulating factor (MSP). This sequence is commodified by the addition of an N-terminal sequence encoding a secretory leader, an initiating methionine proceeding exon II and a terminating codon following exon IV. The cDNA sequence given contains sequences coderived from human megakaryocyte colony stimulating factor (meg-CSF). Commodified by the initiating methionine, and encodes a classical commodified meg-CSF includes exons II-IV and is thought to terminate in the cregion between amino acid residues 11-IV and is thought to terminate in the cregion between amino acid residues 134 - 147. The primary transcript of this gene may be cleaved in different ways to yield a family of mRNA's capablity, folding and processing of the molecule. These exons are also thought to play a role in the observed synergy of MSF with other cytokines. Exons V - XII are believed to be implicated in the processing of folding of these exons may contain sequences which direct proteolytic cormore of these exons may contain sequences which direct proteolytic cheavage, adhesion, organisation of the cellular matrix or extracellular matrix processing. Both naturally occuring and non-naturally occuring conditions of alternatively occuring the correct PN felicle.
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Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human MSF; megakaryocyte stimulating factor; tribonectin; alternative splicing; joint boundary lubricant; O-linked oligosaccharide; osteoarthritis; tribosupplementation; tissue adhesion inhibition; friction coefficient reduction; gene therapy; antiarthritic; osteopathic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human megakaryocyte stimulating factor (MSF), SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB29773 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1404 AA;
Novel tribonectin polypeptide useful as lubricant for treating osteoarthritis, comprises O-linked lubricating moiety.
                                                                                                    WPI; 2001-024673/03.
N-PSDB; AAC81498.
                                                                                                                                                                                                                                                                                                                                                                                                    24-APR-2000; 2000WO-US010953
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                                                                                                                                                                                                                                                                       (RHOD-)
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                                                                                                                                                                                                                                                                       RHODE
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                                                                                                                                                                                                                                                                       ISLAND HOSPITAL LIFESPAN PARTNER
                                                                                                                                                                                                                                                                                                                                     99US-00298970.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 1404 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            86.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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1.5e+03;
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ARBSULT 25
AAB60568
ID AAB60
XX AAB60
XX AAB60
XX 27-AP
XX Human
XX Human
XX Homo
XX OACO
PN WO200
XX 01-FE
XX 21-JU
PR 19-JU
PR 19-JU
PR 23-JU
PR 23-JU
PR 23-JU
PR 19-JU
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         indicates the presence of or predisposition to developing osteoarthritis. The tribonectin and DNA encoding it are useful in the treatment of osteoarthritis, where they may be used for lubricating mammalian joints, such as articulating joints of humans, dogs or horses. The tribonectin, when formulated as a membrane, foam, gel or fibre, is useful for inhibiting adhesion between two surfaces such as the injured tissues of a mammal, where the injury is caused by a surgical insertion or trauma, or an artificial device e.g., an orthopaedic implant. In particular, one of the surfaces is pericardial tissue. DNA encoding a tribonectin may be used in gene therapy. The present sequence represents human MSF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; CACP protein; camptodactyly-arthropathy-coxa vara-pericarditis; MSF; megakaryocyte stimulating factor; synovial lubricant; chromosome 1q25-31; osteoarthritis; joint lubrication; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human megakaryocyte stimulating factor (MSF, CACP).
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                                                            New composition comprising the camptodactyly-arthropathy-coxa varapericarditis protein in combination with an anesthetic, useful for treating osteoarthritis, or as lubricants of tissue and joints.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1404 AA;
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                                                                                                                                                                                                           WPI; 2001-182721/18.
                                                                                                                                                                                                                                                                                                                                                                                                            23-JUL-1999; 99US-0145328P
19-JUL-2000; 2000US-00145328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUL-2000; 2000WO-US020002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-APR-2001
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                                                                                                                                                                                                                                                                        Warman ML;
                                                                                                                                                                                                                                                                                                                                            (UYCA-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                               CASE WESTERN RESERVE
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Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Example 1; Page; 34pp; English

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ADK65839
ID ADK65839
ID ADK65839
ID ADK65839
AC ADK65
XX ADK65
XX ADK65
XX Cytog
KW Cytog
KW angic
KW angic
KW Gene
KW Coror
XX Homo
OS Homo
OS Homo
OS Homo
OS 114-AI
PF 07-FI
PR 10-JI
PR 10-JI
PR 10-JI
PR 03-JI
XX CORI
XX WPI;
XX UDete
PT diag
PT diag
PT diag
PT diag
PT disch
PS Disc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method of treating osteoarthritis via the cambinistration of a composition comprising the camptodactyly-arthropathycox awar-pericarditis (CACP) protein, or portions of the AACP protein.

CC oxa wara-pericarditis (CACP) protein, or portions of the AACP protein.

CC of the invention may further comprise a local anaesthetic. The composition of the invention may be administered via intra-articular or intravenous injection. The human CACP protein is identified in the invention as being megakaryocyte stimulating factor (MSF). The gene encoding CACP protein (MSF) is located on chromosome 1q25-31, and mutations in this gene are cresponsible for the heritable disorder camptodactyly- arthropathy-coxa cox vara-pericarditis, in which patients have synovial hyperplasia without evidence of inflammation. CACP protein (MSF) acts as a synovium comprised to lubricate tissue and joints in the creating factor (MSF). The composition may be applied to reduce the symptoms of osteoarthritis (e.g., joint pain, loss of range of movement cor joint damage). The present sequence represents human megakaryocyte stimulating factor (MSF). CACP protein). Note: This sequence is not given in isometry in figure 4 of the specification, although a GenBank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                      Determining the angiogenic index of a tissue or cell sample using expression levels of differentially expressed genes, useful for diagnosing or treating cancer, coronary artery disease, myocardial ischemia and/or arteriosclerosis.
                                                                                                                                                                                                                                                                    07-FEB-2002; 2002US-00067482:
10-JUN-2002; 2002US-00164595:
16-AUG-2002; 2002US-0403649F:
03-JAN-2003; 2003US-0437746F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      angiogenesis inhibitor; angiogenesis stimulator; angiogenic index; gene expression; cancer; coronary artery disease; myocardial ischemia; coronary arteriosclerosis; forensic medicine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; cardiant; vasotropic; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1404 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       accession number was
                                                                                                                                                                                                                                                                                                                                                                   07-FEB-2003; 2003WO-US003848
                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003066831-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Angiogenesis-differentially expressed protein #53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADK65839 standard; protein; 1404 AA.
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nes 6; Conserv
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                                                                                                                                                                                                                                  ORIGENE TECHNOLOGIES INC
                                                                                                                                                                                         Li X,
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                                                                                                                                                                                           Kovacs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       given. This sequence was therefore obtained
                    78;
                296pp; English
                                                                                                                                                                                           Fan W,
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Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                           Jay
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RESULT 27
ADM9801.4
ID ADM9
XX ADM9
AC ADM9
XX ADM9
XX Lubi
XX Homc
XX US2(
XX
Query Match
Best Local Similarity
Thes 6; Conserve
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Best Local
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                                                                                                                                                                                                                                                      The invention relates to a lubricating polypeptide and at least one O-linked oligosaccharide. The composition and methods are useful for lubricating joints or other tissues to prevent or treat camprodactylarthropathy pericarditis (CAR) or osteoarthritis in mammals. The present sequence represents the amino acid sequence of the human megakaryocyte
                                                                                                                                                            Sequence
                                                                                                                                                                                                                   sequence represents the amino acid sequence of stimulating factor (\mathsf{MSF}).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New tribonectin polypeptides and polynucleotides for lubricating joints or other tissues to prevent or treat Camptodactyl-arthropathy-pericarditis syndrome or osteoarthritis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-APR-1999; 99US-00298970.
24-APR-2000; 2000US-00556246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lubricating polypeptide; O-linked oligosaccharide; joint lubrication;
CAP; camptodactyl-arthropathy pericarditis; osteoarthritis; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-373948/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-JUL-2001; 2001US-00897188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    megakaryocyte stimulating factor; MSF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM98014 standard; protein; 1404 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        megakaryocyte stimulating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JAY G
                                                                                                                                                            1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1; 34pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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Conservative

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86.8%;

Score 33; DB Pred. No. 1.5e 0; Mismatches

; DB 8; . 1.5e+03;

Length 1404;

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Gaps

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RESULT 28
ADX69335
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                                                              Query Match
Best Local S
Matches 6
                                                                                                                                                      The invention relates to heparin-binding vascular endothelial growth factor receptor 3 (VEGFR-3) proteins and encoding polynucleotides. The heparin binding VEGFR-3) proteins are used for stimulating the lymphangiogenesis or angiogenesis in a mammal, and for modulating the growth of mammalian endothelial cells, mammalian endothelial precursor cells or hematopoletic progenitor cells. The polypeptide may also be used for promoting recruitment, proliferation, differentiation, migration or survival of neuronal cells or neuronal precursor cells, and for treating neurodegenerative discorder, e.g. Alzheimer's diseases, Parkinson's disease, Huntington's disease, motor neuron disease, Parkinson's clerosis (ALS), dementia, or cerebral palsy. The present sequence represents a human heparin binding protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heparin binding protein; Neuroprotective; Nootropic; Antiparkinsonian; Anticonvulsant; VEGF-3 receptor; Anglogenesis stimulator; Gene Therapy; vascular endothelial growth factor receptor 3; VEGFR-3; angiogenesis disorder; neurodegenerative disorder; Alzheimers disease; parkinsons disease; motor neurone disease; dementia; paralysis; VEGF-C; neurological disease; Huntingtons chorea;
                                                                                                                                                                                                                                                                                                                                                                                                      comprising prepro-VEGF-C, preating neurodegenerative Huntington's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                     New heparin-binding Vascular Endothelial Growth Factor Receptor 3 ligands comprising prepro-VEGF-C, prepro-VEGF-D sequences or fragments, for treating neurodegenerative disorder, e.g. Alzhelmer's, Parkinson's, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JUN-2003; 2003US-0478114P.
12-JUN-2003; 2003US-0478390P.
23-SEP-2003; 2003US-00669176.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human heparin binding protein #35.
                                                                                                                             Sequence 1404 AA;
                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JUN-2004; 2004WO-US019122
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                                                              Similarity 6; Conserv
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 WMLSPFS 1177
                                WMLSAFS
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                                                                Conservative
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                                                                             86.8%;
                                                                                                                                                                                                                                                                                                                                                                      219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    factor receptor 3; neurological disease
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                                                                            Score 33; DB 9;
Pred. No. 1.5e+03;
                                                                Mismatches
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                                                                                             Length 1404;
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RESULT 29
ADV15658
ID ADV15
XX ADV16
XX
RESULT 30
AED66477
ID AED66677
AC AED66
XX AED66
XX AED66
XX Human
XX Wound
XX Wound
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XX Homo
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated nucleic acid encoding a PR polypeptide. The polypeptide, agonist or an antagonist, antibod composition, and method are useful for diagnosing and treating related disorder, e.g. systemic lupus erythematosus, rheumatoid arthitis. The present sequence represents a DNA encoding a PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New nucleic acid encoding PRO polypeptide, useful for diagnosing and treating an immune related disorder, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, thyroiditis, or diabetes mellitus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antirheumatic; Antiarthritic; Osteopathic; Hemostatic; Antianemic; Antithyroid; Antidiabetic; Nephrotropic; CNS-Gen.; Hepatotropic; Virucide; Gastrointestinal-Gen.; Antipsoriatic; Antiasthmatic;
                                                                                                                                                     Wound healing
                                                                                                                                                                                                   Human lubricin protein sequence.
                                                                                                                                                                                                                                                         29-DEC-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1404 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-AUG-2004; 2004WO-US026249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRO polypeptide SEQ ID NO 1464.
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                                                   Homo sapiens
                                                                                                       vulnerary; injury; lubricin.
                                                                                                                                                                                                                                                                                                                                                                AED66477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1464; 158pp; English.
                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WMLSPFS 1177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                         (first
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                                                                                                                               agent; tissue regeneration; antisense therapy;
jury; fractures; osteopathic; synovitis; inflamm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immune
                                                                                                                                                                                                                                                                                                                                                             protein; 1404 AA
                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouyang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            86.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Williams
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1404;
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RESULT 31
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The new invention relates to the healing and/or repair of cartilaginous tissue. Described is the use of a compound in the preparation of a medicament for promoting the healing or integration of the cartilaginous tissue in a mammal. Given is the use of a compound that inhibits the post translational glycosylation of lubricin in a cell, or a compound having an antisense first nucleic acid sequence to inhibit the synthesis of lubricin in a mammalian cell, where the first nucleic acid sequence is complementary to a fragment of a second nucleic acid sequence, or one substantially identical to it, that encodes lubricin, thus reducing the concentration of lubricin in the extracellular matrix (ECM) that contacts cartilaginous tissue, in the preparation of a medicament for promoting the healing or integration of the cartilaginous tissue in a mammal. The compound inhibits for example, a glycosyltransferase enzyme, e.g. N-acetylgalactosaminyltransferase, or castilaginous tissue, in the preparation of a medicament for promoting the healing or integration of the cartilaginous tissue in a mammal. The compound inhibits for example, a glycosyltransferase enzyme, e.g. N-acetylgalactosaminyltransferase, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     galactosyltransferase. The nucleic acid sequence encoding lubricin comprises SEQ ID NO: 1. The compound is useful or treating blunt trauma, chondral fracture, osteochondral fracture, traumatic synovitis, or damage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Use of a compound, in preparing a medicament for promoting the healing integration of the cartilaginous tissue in a mammal.
                                23-JUL-2004; 2004US-0590766P
                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                     arthropathy; musculoskeletal disease; pharmaceutical; therapeutic
                                                                                                                                                                                                                                     Megakaryocyte stimulating factor (MSF).
                                                                                                                                                                                                                                                                           23-MAR-2006
                                                                                                                                                                                                                                                                                                             AEF54256
                                                                                                                                                                                                                                                                                                                                            AEF54256 standard; protein; 1404 AA.
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                                                                22-JUL-2005; 2005WO-US026004
                                                                                                                                    WO2006012492-A2
 (MUCO-)
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MUCOSAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                           (first entry)
THERAPEUTICS LLC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.8%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or ligaments. The preesnt sequence is human lubricin
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Pred. No. 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  also described is the use of tribonectin in the preparation of a medicament for the lubrication and chondroprotection of a mammalian joint, where the tribonectin is added to a visco-supplement in the preparation of the medicament, which increases the elasticity of the visco-supplement. (A) is useful in the preparation of medicament for lubrication or chondroprotection of a mammalian joint (particularly articulating a joint of a human, horse or dog). The present sequence represents human megakaryocyte stimulating factor (MSF).
                                                                                                                                                         New isolated tribonectin comprising a boundary-lubricating amount polypeptide, and an O-linked oligosaccharide group, for use in trearthritic diseases, and to coat artificial limbs and joints.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Visco-supplementation chondroprotection of rtribonectin.
                                                                                                         Claim 1; SEQ ID NO 1; 34pp; English.
                                                                                                                                                                                                                                                                        WPI; 2006-170699/18.
N-PSDB; AEF89840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-2000; 2000US-00556246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tribonectin; arthritis; antiarthritic; megakaryocyte stimulating factor; MSF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human megakaryocyte stimulating factor (MSF) - SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AEF89839 standard; protein; 1404 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US7001881-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprises hyaluronic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                 (RHOD-) RHODE ISLAND HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1171 WMLSPFS 1177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AEF54257
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00298970
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a viscosupplementation composition (7 (1-5 mg/ml) and tribonectin (10-250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50pp; English.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 10;
1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      osteoarthritis; osteopathic;
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                                                                                                                                                                                         in treating
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The invention comprises a tribonectin protein (a boundary lubricating polypeptide), the tribonectin protein of the invention comprises the human megakaryocyte stimulating factor (MSF) protein. The tribonectin

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RESULT 33
AAU32262
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                polypeptides and antibodies to the polypeptides are useful for identifying agents (agent to account to a polypeptide). The polypeptides are also useful for identifying agents (agentsts and antagonists) that bind to them. Cells expressing the proteins are useful for identifying agents (agentsts and antagonists) that bind to them. Cells expressing the proteins are useful for identifying a therapeutic agent for use in treatment of a pathology related to abservant expression or physiological interactions of the polypeptide. Vectors comprising the nucleic acids encoding the polypeptides and cells genetically engineered to express them are also useful for producing the proteins. The proteins are useful in genetic vaccination, testing and therapy, and can be used as nutritional supplements. They may be used to increase stem cell proliferation; to regulate haematopoiesis; and in bone, cartilage, tendon and/or nerve tissue growth or regeneration; immune suppression and/or stimulation; as anti-inflammatory agents; and in treatment of leukaemias.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein of the invention is useful for treating arthritic disease and osteoarthritic disease, and for coating artificial limbs and joints prior to implantation. The present amino acid sequence represents the human MSF protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1404 AA;
                                                                                                                                                                                                                                                                                                   The invention relates to novel human secreted polypeptides. The polypeptides and antibodies to the polypeptides are useful for
                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-APR-2000; 2000US-00552929.
26-JAN-2001; 2001US-00770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200179449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human secreted protein #2753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAU32262;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAU32262 standard; protein;
Sequence 1415 AA;
                                   AAU29510-AAU33304 represent the amino acid sequences secreted proteins of the invention
                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-611725/70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tang YT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HYSE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-APR-2001; 2001WO-US008656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1171 WMLSPFS 1177
                                                                                                                                                                                                                                                                                                                                                       20; Page 573; 765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Liu C, Drmanac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 33; DB
Pred. No. 1.5e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1404;
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                                                   of novel human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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Query Match

86.8%;

Score 33;

DB 4;

Length 1415;

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RESULT 34
AAM82730
ID AAM82
XX AAM82
XX AAM82
XX AAM82
XX Human
XX Human
XX Human
XX Human
XX Homo
XX 
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Matches 6; Conserv
                                                             18-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
30-AUG-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
                                                                                                                                                                                                                                                                                       14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
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07-JUL-2000;
11-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2000;
14-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JUL-2000
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14-JUL-2000;
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19-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAM82730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAM82730 standard; protein; 31 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1182 WMLSPFS 1188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>س</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; vaccine; metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                     2000US-0217496P
2000US-0220964P
2000US-0220964P
2000US-0224519P
2000US-0225214P
2000US-0225214P
2000US-0225266P
2000US-0225267P
2000US-0225267P
2000US-0225757P
2000US-0225757P
2000US-0225759P
2000US-02259345P
2000US-02259345P
2000US-02259345P
2000US-02259513P
2000US-02259513P
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2000US-0216880P.
2000US-0217487P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000US-0209467P.
2000US-0214886P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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2000US-0231242P.
2000US-02312443P.
2000US-02312413P.
2000US-0231414P.
2000US-0232081P.
2000US-0232396P.
2000US-0232396P.
2000US-0232396P.
2000US-0232396P.
2000US-0232396P.
2000US-0232396P.
2000US-02323065P.
2000US-0233065P.
2000US-0233065P.
2000US-0234074P.
2000US-0234274P.
2000US-0234274P.
2000US-0234274P.
2000US-0234274P.
2000US-0234274P.

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RESULT 35
ABP04103
ID ABP04
XX
XX
AC ABP04
AC ABP04
XX
XX
DT 25-JI
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DT 25-JI
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XX
Humar
XW Humar
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                                                                                                                                                                                                                                                                                                                                                    CC AAKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) CC amino acid sequences given in AAM82170 to AAM93921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CI treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CI that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC nucleic acids into a host cell and culturing the cell to express the CC concers and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK64902 to AAK87694 represent numan immune/haematopoietic antigen genomic CC represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-NOV-2000
17-NOV-2000
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17-NOV-2000
01-DEC-2000
01-DEC-2000
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08-DEC-2000
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08-DEC-2000
08-DEC-2000
08-DEC-2000
Human; open reading frame; ORFX; gene therapy; cancer; cirrhosis; hyperproliferative disorder; psoriasis; benign tumour; haemorrhage;
                                           Human ORFX protein sequence SEQ ID NO:8188.
                                                                                                                                  ABP04103
                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding useful for preventing,
                                                                        25-JUN-2002
                                                                                                       ABP04103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN
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                                                                                                                                                                                                         1 WMLSAFS
                                                                                                                                                                                                                                                                  Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAK55511.
                                                                                                                                 standard; protein; 81
                                                                                                                                                                                                                                                                                                                              31 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 10323; 3071pp +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preventing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0249245P,
2000US-249265P,
2000US-0249297P,
2000US-0249299P,
2000US-02503160P,
2000US-0251030P,
2000US-0251130P,
2000US-0251719P,
2000US-0251866P,
2000US-0251866P,
2000US-0251869P,
2000US-0251989P,
                                                                                                                                                                                                                                                                     Conservative
                                                                        (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SC,
                                                                        entry)
                                                                                                                                                                                                                                                                                 84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human immune/hematopoietic diagnosing and/or treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INC
                                                                                                                                                                                                                                                                    Score 32; DB Pred. No. 39; 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence Listing;
                                                                                                                                                                                                                                                                                 39;
                                                                                                                                                                                                                                                                                               4.
                                                                                                                                                                                                                                                                                               Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antigen polypeptides, cancers and metastasis.
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                                                                                                                                                                                                                                                                    Gaps
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29-SEP-2000
20-OCT-2000
21-NOV-2000

2000US-023498P 2000US-0235834P 2000US-0235836P 2000US-023636P 2000US-023636P 2000US-023636P 2000US-0236370P 2000US-0237039P 2000US-0237039P 2000US-0237039P 2000US-0237039P 2000US-023993FP 2000US-0241251P 2000US-0241251P 2000US-0241786P 2000US-0241786P 2000US-0241786P 2000US-0241786P 2000US-0244617P 2000US-0246474P 2000US-0246474P 2000US-0246474P 2000US-0246474P 2000US-0246479P 2000US-0246479P 2000US-0246479P 2000US-0246523P 2000US-0246523P 2000US-0246513P 2000US-0246513P 2000US-0246513P 2000US-0246513P 2000US-0249211P 2000US-0249211P

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RESULT 36
ADC95262
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                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    degenerative disorder; osteoarthritis; neurodegenerative disorder; cardiovascular disease; diabetes mellitus; systemic lupus erythematosus; hypertension; hypothyroidism; cholesterol ester storage disease; immune deficiency; immune disorder; infectious disease; autoimmune disorder; rheumatoid arthritis; autoimmune thyroiditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel human polypeptides and polynucleotides useful for diagnosing, preventing and treating cardiovascular disease, neurodegenerative, hyperproliferative disorders and autoimmune disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-106308/14.
N-PSDB; ABN19855.
                                                                                                                                                                                                                                                                                                                              Sequence 81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAY-2000; 2000US-0206132P.
29-AUG-2000; 2000US-0228716P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2001; 2001WO-US010836.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   myasthenia gravis.
                   ADC95262;
                                                        ADC95262 standard; protein; 103 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                     format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CURA-) CURAGEN CORP.
                                                                                                                                                               83
                                                                                                                                                                                                  1 WMLSAFS 7
                                                                                                                                                                                                                                         Similarity 5; Conserv
                                                                                                                                                             WLLSAYS 74
                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                              AA:
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                                                                                                                                                                                                                                           Score 32; DB 5;
Pred. No. 1.1e+02;
2; Mismatches 0
                                                                                                                                                                                                                                                                                  Length 81
                                                                                                                                                                                                                                             Indels
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AAU25590
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                                                                                                                                                    RESULT 37
                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                    recombinant vector comprising the nucleic acid operably linked to transcription regulatory element, a cell comprising the vector and a single-stranded probe comprising the nucleic acid. The nucleic acids are chosen from 3654 disclosed sequences encoding 3654 disclosed proteins. The nucleic acids is useful for diagnosing pathological conditions resulting from E. faecium bacterial infection (e.g. urinary tract infection, bacteraemia, endocarditis, wounds and abdominal-pelvic infection) and for screening drugs such as agonists and antagonists. The nucleic acid is useful for recombinant production of Candida albicans derived peptides or antisense polypeptides. Pharmaceutical compositions and vaccines containing the nucleic acid are useful for preventing or treating Enterococcus faecium infections. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vaccine; urinary tract infection; bacteraemia; endocarditis; wound; abdominal-pelvic infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Enterococcus faecium encoding an Enterococcus faecium polypeptide having one of 10 fully defined sequences given in the (or comprising 40 sequential nucleotides chosen from any of the nucleic acids, its complement or sequences hybridising to it). Also included are a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated nucleic acid derived from Enterococcus faecium encoding Enterococcus faecium polypeptide useful for detection, prevention and treatment of a pathological condition resulting from a bacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6583275-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enterococcus
                                                             18-DEC-2001
                                                                                                                      AAU25590 standard; protein;
                                                                                                                                                                                                                                                                                                                          Sequence 103 AA;
                                                                                                                                                                                                                                                                                                                                                       treating Enterococcus faecium infections. one if the disclosed E. faecium proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; SEQ ID NO 4889;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Doucette-Stamm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENO-) GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 faecium protein sequence SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003-799836/75.
DB; ADC91608.
                                                                                                                                                                                                  17
                                                                                                                                                                                                                                                          Similarity
5; Conserv
                                                                                                                                                                                                                               1 WMLSAF 6
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WMISAF
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                                                             (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0051571P.
98US-0085598P.
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                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bush D;
                                                                                                                                                                                                                                                                            84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         243pp; English
                                                                                                                                                                                                                                                            Score 32; DB
Pred. No. 1.4e
1; Mismatches
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                                                                                                                                                                                                                                                         Db
1.4e+02;
0;
                                                                                                                                                                                                                                                                                            Length 103;
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                                                                                                                                                                                                                                                                                                                                                                          represents
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Human; G-protein coupled receptor; GPCR; mental disorder; schizophrenia;

Human G Protein-Coupled Receptor (GPCR) polypeptide #37.

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Matches
                                                                                                                                 Query Match
Best Local Similarity
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23-FEB-2000;
02-MAR-2000;
03-MAR-2000;
09-MAR-2000;
13-MAR-2000;
                                                                                                                                                                                                                                                                                                          Sequences AAU25554-AAU25616 represent human G-protein coupled receptor (GPCR) polypeptides of the invention. The proteins and their associated DNA sequences can be used to identify compounds which bind to GPCR polypeptides and in screening for compounds that modulate GPCR activity. By screening a human subject for the presence of mutations in GPCR DNA, a GPCR-related disorder or a genetic predisposition can be diagnosed. The sequences can also be used for treatment and prevention of mental disorders such as schizophrenia, attention deficit disorder, anxiety, depression, dementia and bipolar disorder, neurological disorders such as Huntington's disease, Parkinson's disease and Tourette's syndrome, metabolic disorders such as obesity, anorexia and type 2 diabetes, cardiowascular disorders such as thrombosis, myocardial infarction, and atheroscolarsis with infarctions canced by HIV and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid encoding a new G-protein coupled receptor polypeptide for detecting receptor modulators that can treat mental disorders, such as schizophrenia, anxiety, depression, or obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    viral infection; immunostimulant; neuroleptic; nootropic; tranquiliser;
antidepressant; anorectic; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 35;
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                                                                                                                                                                                                                Sequence 211
                                                                                                                                                                                                                                                                                               cardiomyopathy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  attention deficit disorder; anxiety; depression; bipolar disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA & UPJOHN CO
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128
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WLLAAFS 134
                                                     WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0184247P.
2000US-0184303P.
2000US-0184305P.
2000US-0184397P.
2000US-0186457P.
2000US-0186810P.
2000US-018681P.
2000US-018880P.
2000US-013861P.
2000US-0213861P.
2000US-02137369P.
2000US-021837P.
2000US-021837P.
2000US-021837P.
                                                                                                        Conservative
                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                               and atherosclerosis, viral infections caused by HIV and
                                                                                                                              84.2%;
71.4%;
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                                                                                                                                 Score 32; DB 4;
Pred. No. 3.1e+02
                                                                                                        Mismatches
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                                                                                                     Gaps
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RESULT 38 ADR09699

RESULT 39
ABO73613
ID ABO73

AB073613

standard; protein; 262

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Best Local &
                                                                                                                                                                                                                                                                                                                                                             This invention relates to novel, isolated full length human cDNA molecules and the encoded proteins thereof. Specifically, it refers to cDNA clones obtained by an oligo-capping method, where none of these clones are identical to any known human mRNAs. The present invention describes an immunoassay to identify agonists and antagonists, as well as antibodies, antisense molecules and sink that can all be used to bind to and modulate expression of the cDNA molecules. As such, these molecules are useful for diagnostic markers or therapeutic targets for
                                                                                                                                                                                     the various diseases or morbid states. In particular, they are useful in gene therapy for treating osteoporosis, neurological disease, Alzheimer's disease, Parkinson's disease, dementia, short memory and various cancers, as well as for maintaining equilibrium of sense or motor function, and for treating emotional reaction, fear response and panic. Accordingly, they exhibit osteopathic, neuroprotective, noutropic, antiparkinsonian, cytostatic and tranquiliser activities. This polypeptide is a protein encoded by a full length human cDNA sequence of the invention. NOTE: This sequence is not given in the sequence listing of the specification but can be obtained on CD-ROM from the European Patent Office, Vienna Sub-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             osteoporosis; neurological disease; Alzheimer's disease;
Parkinson's disease; dementia; short memory; cancer;
sense or motor function; emotional reaction; fear response; panic;
osteopathic; neuroprotective; nootropic; antiparkinsonian; cytostatic;
                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New 1995 cDNA, useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-583265/57.
N-PSDB; ADR07743.
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Wakamatsu A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-FEB-2003; 2003JP-00102207.
09-MAY-2003; 2003JP-00131452.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New 1995 cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-FEB-2004; 2004EP-00003145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tranquiliser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REAS-) RES ASSOC BIOTECHNOLOGY
 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein useful for treating neurological disease Seq 3205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligo-capping method; diagnostic marker; gene the orosis; neurological disease; Alzheimer's disease;
                                1 WMLSAFS 7
                                                                                   Similarity
                                                                  6
WKLSAFS 193
                                                                                                                                     243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yamamoto J,
A, Ishii S,
                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NO 3205;
                                                                                   84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nagai K, Irie R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2686pp; English.
                                                                  0
                                                                  Score 32; DB Pred. No. 3.6e 0; Mismatches
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                                                                                   .6e+02;
                                                                                                   Length 243;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy;
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RESULT 40
AAG81234
ID AAG81
XX
AC AAG81
XX
DT 04-SE
XX
DE Mycok
XX
KW Drug
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Best Local Similarity
                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                  including anti-P. aeruginosa drugs, as templates for recombinant production of P. aeruginosa-derived peptides or polypeptides, as target components for diagnosis and/or treatment of P. aeruginosa-caused infection, and in detection of P. aeruginosa sequences or other sequences of Pseudomonas species using biochip technology. Sequences ABO67826-ABO84396 represent P. aeruginosa polypetides of the printed sequence data for this patent did not form part of the printed specification but was obtained in electronic format from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to Pseudomonas aeruginosa polypeptides and the polynucleotides encoding them. The sequences are useful in diagnosis and therapy of pathological conditions, as molecular targets for diagnostics, prophylaxis and treatment of pathological conditions resulting from a bacterial infection, for evaluating a compound, such as a polypeptide, for the ability to bind a P. aeruginosa nucleic acid, as components of effective antibacterial targets, as targets for antibacterial drugs,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated nucleic acid encoding Pseudomonas aeruginosa polypeptide, useful as molecular targets for diagnostics, prophylaxis and treatment c pathological conditions resulting from bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rubenfield MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-1998;
27-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-APR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pseudomonas aeruginosa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacterial infection; Pseudomonas aeruginosa infection; antibacterial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pseudomonas aeruginosa polypeptide #5788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JUL-2004 (first entry)
             Drug target; growth; organism viability; characterisation
                                               Mycobacterium tuberculosis potential drug target protein SEQ
                                                                             04-SEP-2001
                                                                                                                                                                                                                                                                                                                                                      Sequence 262 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 22359; 455pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO73613;
                                                                                                                                          AAG81234 standard; protein; 280 AA
                                                                                                                                                                                                                                                                                                                                                                                   seqdata.uspto.gov/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENO-) GENOME
                                                                                                             AAG81234;
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                                                                                                                                                                                                                         234 WMLSAIS 240
                                                                                                                                                                                                                                                       1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                         Conservative
                                                                           (first entry)
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98US-0094190P.
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                                                                                                                                                                                                                                                                                                       84.2%;
                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                       Score 32; DB 7;
Pred. No. 3.9e+02
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RESULT 41
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide sequence that may be a drug target, or essential for growth or viability of an organism. Polymucleotide sequences AH51947 - AH52092 represent DNA encoding proteins AAG81096 - AAG81241, Mycobacterium tuberculosis proteins which are potential drug targets. The DNA and protein sequences are used to illustrate the method of the invention. The method involves providing an unknown nucleotide or polypeptide sequences, and comparing it to a number of sequences along with at least one algorithm capable of analysing a functional relationship between nucleotide and polypeptide sequences. The method is useful for characterising the function of nucleic acids and polypeptides that may be useful as a target for a drug or essential for the growth or viability of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-NOV-1999; 99US-0165086P
12-NOV-1999; 99US-0165124P
01-FEB-2000; 2000US-0179531P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying nucleotide or polypeptide sequence for use as drug target, involves providing algorithm that analyzes a functional relationship between nucleotide or polypeptide sequences, and comparing the sequences.
24-APR-2000; 2000US-0199149P.
04-AUG-2000; 2000US-00633146.
                                                                                       24-APR-2001; 2001WO-US013097
                                                                                                                                                  01-NOV-2001
                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                Human GPCR polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAM51647;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAM51647 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 280 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  an organism
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 190; 207pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-NOV-2000; 2000WO-US031152
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                                                                                                                                                                                                             WO200181409-A2
                                                                                                                                                                                                                                                                                                                                                         Human; G-protein coupled receptor; GPCR; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (REGC ) UNIV
                                                                                                                                                                                                                                                                                                                         proto-oncogene receptor; human protease; disease
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71.4%;
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Pred. No. 4.2e+02;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated human G protein-coupled receptor (GPCR) polypeptide that is related to the MAS prote-oncogene receptor subfamily. The polypeptide comprises a fully defined sequence of 289 amino acids as given in the specification, or its fragment comprising 10 contiguous amino acids, or an amino acid sequence of an allelic variant or orthologue of the amino acid sequence given in the sequence. The polypeptide is useful for identifying a modulator of a GPCR polypeptide or an an agent that binds to it. The polypeptide is also useful for treating a disease or condition mediated by human proteases. The present sequence is the polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                          G protein-coupled receptor; GPCR; drug screening; diagnosis; transgenic mouse; neurological disorder; adrenal gland disorder; colon disorder; intestinal disorder; cardiovascular disorder; muscular disorder; blood disorder; immune disorder; blood disorder; immune disorder; cancer; kidney disorder; metabolic disorder; nutritive disorder; cancer; kidney disorder; uterus disorder; lung disorder; treast disorder; ovary disorder; uterus disorder; prostate disorder; testis disorder; skin disorder; stomach disorder; pancreas disorder; spleen disorder; thymus disorder; thyroid disorder; antiparkinsonian; antimanic; cytostatic; antiflammatory; vasotropic; antidiarrhoeic; antidiabetic; virucide; hepatotropic; antibacterial; antianaemic; antiseborrhoeic; dermatological; antibleer; antithyroid; antiallergic; anorectic;
                               09-SEP-2002; 2002US-0409303P
09-APR-2003; 2003US-0461329P
                                                                                                               09-SEP-2003; 2003WO-US028226.
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                                                                                                                                                                                13-MAY-2004.
                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mouse GPCR
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                                                                                                                                                                                                                                                                                                                                                                                 immunosuppressive; nephrotropic; gene therapy; GPCR modulator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PEKE )
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DB; ABA03701.
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                                                                                                                                                                                                                                                                                                                                                        receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MRGG), SEQ ID NO:657.
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Pred. No. 4.3e+02;
2; Mismatches 0
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Human GPCR

N8

(MRGG), SEQ ID NO:656.

(first entry)

G protein-coupled receptor; GPCR; drug screening; diagnosis; transgenic mouse; neurological disorder; adrenal gland disorder; colon disorder; intestinal disorder; cardiovascular disorder; muscular disorder; blood disorder; immune disorder; bone disorder joint disorder; metabolic disorder; nutritive disorder; cancer;

disorder;

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94

WLLAAFS 100

1 WMLSAFS Similarity 5; Conserv

7

Matches Query Match Best Local

Conservative

2

Score 32; DB 8; Pred. No. 4.3e+02; 2; Mismatches 0

<u>,</u>

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0

Length 289;

84.2%; 71.4%;

Sequence

289 AA;

RESULT 43

ADO29554 standard; protein;

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CC (GPCES) and mucleic acids encoding them. The invention also relates to compenses at least 90% identical to the GPCR proteins and nucleic acids cof the invention; methods of treating, preventing or diagnosing diseases comprising a GPCR gene of the invention; methods of screening for compounds useful in the treatment of GPCR related diseases; a transgenic compuse comprising a GPCR gene of the invention; a mouse comprising a GPCR gene or in an endogenous GPCR gene; cells derived from the transgenic mice; kits comprising several mice, each of which has a mutation in a different GPCR pellowing several mice, each of which has a mutation further discloses variants of the GPCR polypeptides and vectors comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may be used in the diagnosis, treatment or prevention of a wide variety of diseases including neurological disorders (e.g., Alzheimer's disease, comparising a GPCR polypeptides and proteins may be used in the diagnosis, treatment or prevention of a wide variety of compression, diabetic neuropathy, Parkinson's disease or schizophrenia); cc disorders of the adrenal gland; disorders (e.g., Alzheimer's disease, comparising of the adrenal gland; disorders of the colon or intestine compocardial infarction); muscular disorders of the colon or intestine compocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or myocardial infarction); muscular disorders; blood disorders (e.g., autoimmune disorders (e.g., osteoarthritis, rheumatoid composity, enzyme deficiency-related diseases or vitamin deficiency-related diseases or vitamin deficiency-related diseases, posteoarthritis, composition of the skin, stomach, pancreas, spleen, thymus and thyroid (e.g., cancers). The present sequences not shown were obtained in celetronic format directly from WIPO at the printed disorder of the colon of the 
Novel mammalian G protein coupled receptors, useful for identifyi compounds that modulates diagnosing and treating disease conditio associated with GPCR dysfunction e.g. autoimmune diseases, angina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 151; SEQ ID NO 657; 542pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates
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lwain KL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 to human and mouse G protein-coupled receptors
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Pavlova MN, Vassilatis D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for identifying ting disease condition
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Zeng H;
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compounds useful in the treatment of GPCR-related diseases; a transgenic compounds useful in the treatment of GPCR-related diseases; a transgenic computed on the transgenic mouse comprising a GPCR gene of the invention; a mouse comprising a crange of the invention; and comprising a crange of the invention; and kits comprising several mice, each of which has a different GPCR gene of the invention; and kits comprising comprising comprising envention further discloses variants of the GPCR polypeptides and vectors comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may be used in the diagnosis, treatment or prevention of a wide variety of diseases including neurological disorders (e.g., Alphaimer's disease, comprising disorders of the colon or intestine comprising crandiovascular disorders of the colon or intestine comprised disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders; bood disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders; bood disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders; bood disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders; bood disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or unyocardiac arrh
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                                                     anaemia or leukaemia); immune disorders (e.g., autoimmune disorders or AIDS); bone and joint disorders (e.g., osteoarthritis, rheumatoid arthritis, gout or osteoporosis); metabolic or nutritive disorders (e.g., obesity, enzyme deficiency-related diseases or vitamin deficiency-related diseases); and disorders of the kidney, liver, lung, breast, ovary, uterus, prostate, testis, skin, stomach, pancreas, spleen, thymus and thyroid (e.g., cancers). The present sequence represents a GPCR of the invention. Note: The full sequence data for this patent did not form part of the printed specification; those sequences not shown were obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel mammalian G protein coupled receptors, useful for identifying compounds that modulates diagnosing and treating disease condition associated with GPCR dysfunction e.g. autoimmune diseases, angina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-SEP-2002; 2002US-0409303P
09-APR-2003; 2003US-0461329P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GPCRs) and nucleic acids encoding them. The invention also relates to sequences at least 90% identical to the GPCR proteins and nucleic acid
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lova MN, Vassilatis
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Query Match Best Local Similarity

84 71

Score Pred.

32; No.

DB 8; 4.3e+02;

Length

Sequences AAU2554-AAU25616 represent human G-protein coupled receptor (GFCR) polypeptides of the invention. The proteins and their associated DNA sequences can be used to identify compounds which bind to GFCR polypeptides and in screening for compounds that modulate GFCR activity. By screening a human subject for the presence of mutations in GFCR DNA, & GFCR-related disorder or a genetic predisposition can be diagnosed. The sequences can also be used for treatment and prevention of mental disorders such as schizophrenia, attention deficit disorder, anxiety, depression, dementia and bipolar disorder, any province a surface and many procession as a compound as a compo

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Matches
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23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
23-FEB-2000;
02-MAR-2000;
09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            attention deficit disorder; anxiety; depression; bipolar disorder; neurological disorder; Huntington's disease; dementia; obesity; anxiety disease; dementia; obesity; anc metabolic disorder; Parkinson's disease; Tourette's syndrome; throm type 2 diabetes; cardiovascular disorder; myocardial infarction; cardiomyopathy; atherosclerosis; human immunodeficiency virus; HIV; viral infection; immunostimulant; neuroleptic.
                                                                                                                             New isolated nucleic acid encoding a new G-protein coupled receptor polypeptide for detecting receptor modulators that can treat mental disorders, such as schizophrenia, anxiety, depression, or obesity.
                                                                                                                                                                                                                                                               11-JUL-2000;
11-JUL-2000;
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23-JUN-2000;
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13-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antidepressant; anorectic; gene therapy.
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2000US-0186457P
2000US-0186457P
2000US-0186810P
2000US-0188880P
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2000US-0218492P
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2000US-0213861P.
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RESULT 45
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Matches 5
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                                                                                                        The present sequence represents a human guanine nucleotide binding protein (G protein)-coupled receptor protein designated TGR15. TGR15 has nootropic, antiinflammatory, vasotropic, immunomodulator and cytostatic activities. The TGR15 polynucleotide and protein can be used in gene therapy and protein therapy. G-protein coupled receptor proteins are cell membrane proteins which mediate the cellular response to a large variety of signalling molecules. The TGR15 polynucleotide and protein can be used in the diagnosis, treatment and prevention of diseases including neurological, inflammatory, circulatory, degenerative, immune system and digestive diseases and cancer
                                                                                                                                                                                                                                                                                      G-protein coupled receptor protein TGR15 of human origin and DNA encoding it for diagnosis and treatment of cancer and circulatory and other diseases associated with its expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            metabolic disorders such as obesity, anorexia and type 2 diabetes, cardiovascular disorders such as thrombosis, myocardial infarction, cardiomyopathy and atherosclerosis, viral infections caused by HIV
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26-DEC-2000; 2000JP-00395566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vasotropic; immunomodulator; cytostatic;
neurological; inflammatory; circulatory;
                                                                                Sequence 323
                                                                                                                                                                                                                                                            Claim 1; Fig 2; 103pp; Japanese
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                                          Local
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DB; ABL40211.
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                            Similarity
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Pred. No. 4.9e+02,
2; Mismatches
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4.9e+02;
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degenerative; immune system;
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16-MAR-2001; 2001US-0276873P.
30-MAR-2001; 2001US-0280531P.
30-MAR-2001; 2001US-0280596P.
16-NOV-2001; 2001US-0334266P.
28-NOV-2001; 2001US-0334229P.
11-JAN-2002; 2002US-0347703P.
                                                                                                                                                                                                   The invention comprises the amino acid and coding sequences of human secreted proteins (SECP). The human SECP DNA and protein sequences of tinvention are useful for the treatment and prevention of cell proliferative disorders (e.g. actinic keratosis, arteriosclerosis, bursitis, hepatitis or cancer); autoimmune/inflammatory disorders (e.g. AIDS, asthma, anaemia, allergies or atopic dermatitis); cardiovascular disorders (e.g. congestive heart failure, ischaemic heart disease, myocardial infarction, hypertensive heart disease, or vascular tumours) neurological disorders (e.g. epilepsy, stroke, cerebral neoplasms, or Alzheimer's disease); and developmental disorders (e.g. renal tubular acidosis, Cushing's syndrome, Duchenne and Becker muscular dystrophy, chypothyroidism). The present amino acid sequence represents a human secreted protein (SECP) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AIDS; asthma; anaemia; allergy; atopic dermatitis; myocardial infarction; cardiovascular disorder; vascular tumour; neurological disorder; stroke; epilepsy; cerebral neeplasm; Alzheimer's disease; developmental disorder; Cushing's syndrome; muscular dystrophy.
                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human secreted proteins and nucleic acids useful in diagnosing, treating and preventing cell proliferative, autoimmune/inflammatory, cardiovascular, neurological, and developmental disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gururajan R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thangavelu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABJ10905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INCY-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |:|:|||
128 WLLAAFS 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yang J,
relu K, G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INCYTE GENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; protein; 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                   B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kallick DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           J, Elliott VS, Duggan BM,
Gietzen KJ, Forsythe IJ,
Lal PG, Baughn MR, Xu Y,
Kallick DA, Walia NK, Mass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         126-127; 162pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mason
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Honchell CD, Lee S;
Lu DAM, Griffin JA;
Tang YT, Azimzai Y;
on PM, Tran UK;
                                                                                                                                                                                                                                                                                                                                                                      tumours);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of the
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Conservative

0;

Score 32; DB 5; Pred. No. 5.7e+02; 0; Mismatches 1

Length 371; Indels

0

Gaps

0

84.2%;

WMLSAFS

7

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RESULT 47
ABO60886
ID ABO60
XX ABO60
XX ABO60
XX Recon
XX Recon
XX Recon
XX Recon
XX Recon
XX Recon
XX Rebs
XX Rebs
XX Rebs
XX Rebs
XX Rescon
XX R
ADF70484
ID ADF7
XX
AC ADF7
XX
AC ADF7
XX
AC 12-:
XX
XX
UT 12-:
XX
UT 12-:
XX
CFP
XX
CGP
XX
CGP
XX
GFP
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       δ
                                                                                                                                                                                                                                                                                                                                                        RESULT 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 6
                         ligand; orphan receptor protein; fusion protein; fluorescent protein; cell expression; green fluorescent protein; GFP; GFP-1; wild-type GFP; GFPuv; Enhanced GFP; EGFP; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes a new isolated nucleic acid encoding a Klebsiel pneumoniae polypeptide. Also described are: a recombinant expression vector comprising the nucleic acid, operably linked to a transcription regulatory element; and a cell comprising the recombinant expression vector. The nucleic acid is useful for preparing a vaccine composition against Klebsiella pneumoniae. This is the amino acid sequence of a Klebsiella pneumoniae polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid encoding a Klebsiella pneumoniae polypeptide, useful for preparing a vaccine composition against Klebsiella pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 7403; 932pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Breton GL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6610836-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Klebsiella pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Klebsiella
                                                                                                                                       Orphan receptor
                                                                                                                                                                                             12-FEB-2004
                                                                                                                                                                                                                                                 ADF70484;
                                                                                                                                                                                                                                                                                                  ADF70484 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JAN-2000; 2000US-00489039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Klebsiella
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO60886;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO60886 standard; protein; 437 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENO-) GENOME THERAPEUTICS CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2003-895346/82.
                                                                                                                                                                                                                                                                                                                                                                                                                                           391 WSLSAFS 397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WKLSAFS 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pneumoniae polypeptide seqid 7403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pneumoniae protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Osborne M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0117747P.
                                                                                                                                       ligand-related human
                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 32; DB
Pred. No. 6.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                    561 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcription regulatory element; antibacterial; Vaccine.
                                                                                                                                    protein SeqID107.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7; Leas, 6.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Klebsiella
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 49
ADU02401
ID ADU02
XX ADU02
XX ADU02
XX ADU02
XX ADU02
XX Cytos
XX Cytos
XX Cytos
XX Droli
XX Droli
XX Droli
XX Homo

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method of identifying ligands to an orphan receptor protein which comprises transforming cells with DNA encoding a fusion protein of the orphan receptor with a fluorescent protein, so that the fusion protein is expressed in the cells (or cell membranes isolated from them) and contacting the cells with the potential ligand to be tested. A suitable fluorescent protein for incorporation in the fusion protein is green fluorescent protein (GFP), for example GFP-1, wild-type GFP, GFPuv or Emhanced GFP (EGFP). The method is useful for the identification of ligands binding to an orphan receptor protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-FEB-2002; 2002JP-00045728
23-JUL-2002; 2002JP-00213949
11-OCT-2002; 2002JP-00298237
  18-APR-2003; 2003US-0463708P.
18-APR-2003; 2003US-0463732P.
02-MAY-2003; 2003US-0467199P.
02-MAY-2003; 2003US-0467230P.
                                                                                                                                                                                                                                                                                                                                                                   cytostatic; antipsoriatic; antiinflammatory; gene therapy; Nanodisc; proliferative disorder; inflammatory disorder; immune disorder; metabolic disorder; bone disorder; CNS disorder; cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 561 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Transformation of cells with a fusion protein protein with a fluorescent protein useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hinuma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2003071272-A1
                                                                                                                                                                                       04-NOV-2004
                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                             ulcerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human polypeptide seqid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JAN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADU02401;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADU02401 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to the orphan receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; ADF70586.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-FEB-2003; 2003WO-JP001901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-AUG-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                  19-APR-2004; 2004WO-US012047
                                                                                                                                                                                                                                         WO2004093804-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-697654/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          128 WLLAAFS 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fujii R,
                                                                                                                                                                                                                                                                                                                                             colitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 1180
                                                                                                                                                                                                                                                                                                                                             human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ogi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 594pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 32;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               868.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8.8e+02;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kawamata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of an orphan receptor identification of ligands
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hosoya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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host cell transformed, transfected, transduced or infected with the concleic acid molecule; a nucleic acid composition composition comprising a carrier or a buffer and one or more compositions comprising the nucleic acid composition for nucleic acid composition one or more compositions comprising the pulpeptide; an animal injected with the polypeptide; a polypeptide composition comprising the polypeptide and a carrier or buffer; a cell culture medium comprising the polypeptide or transfected cells culture medium comprising the polypeptide; making a transformed, transfected, culture medium comprising a bynamic system; ynthesising Manodiscs simultaneously cand for synthesising a dynamic system; yrapting a hydrophobic protein crystal structure; immunising a non-human animal; complement; determining the presence of the nucleic acid molecule or its complement; determining the presence of an antibody to the polypeptide in a sample; an antibody specifically recognising, binding to or modulating call molecule or its biological activity of at least one polypeptide encoded by a nucleic call molecule or its biological critically active fragment; an antibody a nucleic call molecule or its biological with presence of the nucleic acid molecule or its biological with presence of an antibody a nucleic call molecule or its biological activity of at least one polypeptide encoded by a nucleic call molecule or its biological with presence of the nucleic acid molecule or its biological with presence of an antibody and acid molecule or its biological with presence of an antibody specifically recognising.
                                                                   antibody; diagnosing a disease, disorder, syndrome, or condition.

C comprising cancer, or proliferative, inflammatory, immune, metabolic, bone, (NS, genetic, bacterial and viral diseases, disorders, syndromes or conditions in a patient; a modulator composition comprising a modulator and a carrier; gene therapy; prophylactic or therapeutic treatment of a subject; an isolated modified cell comprising at least one first heterologous nucleic acid molecule, where the first heterologous nucleic acid molecule, where the sequence that encodes a first polypuctide; a non-human animal deficient in the polypeptide or that over-expresses the polypeptide; isolated tissues derived from the non-human animal; and one or more cells derived from the non-human animal; and one or more cells derived from the non-human animal. The nucleic acid is useful in preparing a composition for first is the amino acid sequence of a novel human polypeptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUL-2003;
08-JUL-2003;
14-JUL-2003;
14-JUL-2003;
08-AUG-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid molecule or its biologically active fragment; an antibody composition comprising the antibody and a carrier; a bacteriophage, where the antibody is displayed on the bacteriophage; a bacterial cell comprising the bacteriophage; a non-human animal injected with the antibody composition; a host cell that secretes the antibody; making an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes a new first nucleic acid molecule comprising a polynucleotide sequence given in the specification. Also described are: an animal injected with the nucleic acid molecule comprising a second polynucleotide sequence that is at least about 70, 80, 90 or 95% homologous to the first nucleic acid molecule or that hybridises to the first polynucleotide sequence under high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  stringency conditions; a vector comprising the nucleic acid molecule and a promoter that drives the expression of the nucleic acid molecule; a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New first nucleic acid molecule comprising a polynucleotide sequence given in the specification, useful in preparing a composition for diagnosing or treating e.g., cancer, psoriasis or ulcerative colitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 14; SEQ ID NO 868; 291pp; English.
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                                  Query Match
                                                                                                                                                                                                                                                                                                                                                           useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating disorders involving aberrant protein expression or biological activity. The polypeptide and polymucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits to assess biodiversity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reaction (PCR) primers, oligomers, and for chromosome and gene mappi
and in recombinant production of (II). The polymucleotides are also
in diagnostics as expressed sequence tags for identifying expressed
genes. (I) is useful in gene therapy techniques to restore normal
activity of (II) or to treat disease states involving (II). (II) is
                                                                                                                                                   amino acid sequences. ABG00010-ABG30377 represent novel human diagnostic amino acid sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pot_sequences
                                                                                                   Sequence 1923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated polynucleotide and encoded polypeptides, useful diagnostics, forensics, gene mapping, identification of mutal responsible for genetic disorders or other traits and to asse
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RESULT 2 T30536 Typothetical protein 2 - Fugu rubripes (fragment) C.Species: Fugu rubripes C.Species: Fugu rubripes C.Accession: T30536 R.Riboldi Tunnicliffe, G.R.; Platzer, M.; Nyakatura, submitted to the EMBL Data Library, September 1997	Qy 1 WMLSAFS 7 Db 372 WMLSCFS 378	Query Match 89.5%; Score 34; DB; Best Local Similarity 85.7%; Pred. No. 32; Matches 6; Conservative 0; Mismatches	A;Status: preliminary A;Molecule type: DNA A;Residues: 1-637 <gut> A;Cross-references: UNIPARC:UPI00000B56E1; GB:U17417; C;Superfamily: beta-galactosidase, BgaH type C;Keywords: glycosidase; hydrolase</gut>	; Kasmir, J.J.; 995 ne and beta-gala D:95238267; PMID	ALIGNMENTS) isozyme 12 - evision 03-Oct-	468 26 68.4 412 2 G64059 470 26 68.4 417 2 D82172 471 26 68.4 421 2 F83852 472 26 68.4 421 2 F83852 473 26 68.4 421 2 F83852 477 26 68.4 422 2 A55044 477 26 68.4 431 2 T24941 477 26 68.4 435 1 S18609 477 26 68.4 435 1 S18609 478 26 68.4 435 2 JC40441 479 26 68.4 435 2 JC40441 480 26 68.4 430 2 G70693 481 26 68.4 440 2 G70693 482 26 68.4 440 2 G70693 484 26 68.4 444 2 B3238 486 26 68.4 444 2 B3238 487 26 68.4 444 2 B3238 489 26 68.4 447 2 T09414 490 26 68.4 447 2 T09414 491 26 68.4 447 2 T09414 492 26 68.4 450 2 D821373 494 26 68.4 450 2 D821373 496 26 68.4 450 2 D82128 497 26 68.4 450 2 D82128 499 26 68.4 450 2 D82264 500 26 68.4 463 2 T39629 499 26 68.4 463 2 T39629 499 26 68.4 463 2 T39629
:) 100 #text_change 09-Jul-2004 197 . G.; Elgar, G.S.; Brenner, S.; Rosent		2; Length 637;1; Indels 0; Gaps 0;	.7417; NID:g676845; PIDN:AAA75601.1; PID:ç	Brenchley, J.E. ctosidase isozyme from a psychrotrophic A: 1:7721689	Arthrobacter sp. (strain B7) 1995 #text_change 16-Aug-2004	probable serine tr serine transport p serine transporter glutamate dehydrog hypothetical prote beta-4C-adrenergic hypothetical prote ABC transporter (p ammonium transport glutamate dehydrog probable sugar tra regeneration assoc probable dinf prot hypothetical prote protein F14J16.24 transcription fact transcription fact transcription fact transcription fact transcription fact probable Na+/H+ an nickel resistance hypothetical prote pectinesterase hom pmbA protein - Esc permease VCA0904 (maturation of anti transporter homolo hypothetical prote hypothetical prote

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sidase isozyme from a psychrotrophic A:
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 1; Indels
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A;Cross-references: UNIPROT:P72049;
A;Experimental source: strain H37Rv
C;Genetics:
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C70696
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A;Accession: T30536
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Cross-references: UNIPROT:Q8Z0PO; UNIPARC:UPI00000CDBF9; GB:BA000019; PIDN:BAB77574.1
A;Experimental source: strain PCC 7120
C;Genetics:
                                                                                                                                                                                                                                R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, NDNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamenton A;Reference number: AB1807; MUID:21595285; PMID:1179
                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein all0050 [imported] - Nostoc sp. C;Speciles: Nostoc sp. PCC 7120 A;Note: Nostoc sp. strain PCC 7120 is a synonym of Ar C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #c;Pate: 14-Dec-2001 #sequence_revision 14-Dec-2001 #c;Pate: 14-Dec-2001 #sequence_revision 14-Dec-2001 #sequence_revisi
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A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.; A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230
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A;Residues: 1-187 <RIB>
A;Cross-references: UNIPROT:073698; UNIPARC:UPI000001B8E6; EMBL:AF026198; NID:g3098263;
C;Genetics:
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Best Local
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M.; Yamada,
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; Yasuda, M.; Tabata,
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R;Telford, E.A.R.; Watson, M.S.; McBride, K.; Davison, J
Virology 189, 304-316, 1992
A;Title: The DNA sequence of equine herpesvirus-1.
A;Reference number: A41831; MUID:92295566; PMID:1318606
A;Contents: annotation; possible protein-coding frames
A;Note: neither amino acid nor nucleotide sequence is g:
                                                                                                                                                                                                                                                                                                                                                                                                            51.3K capsid protein - equine herpesvirus 1 (strain Ab4p) C;Species: equine herpesvirus 1 C;Species: equine herpesvirus 1 A;Note: host Equus caballus (domestic horse) C;Date: 30-Sep-1992 #sequence_revision 30-Sep-1992 #text_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-92 <STO>
A;Cross-references: UNIPROT:Q9KAL1; UNIPARC:UPI00000C3E6F; GB:AP001515; GB:BA000004; NID
A;Experimental source: strain C-125
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C;Keywords:
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R; Telford, E.A.R.; V
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A; Accession: E36797
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RESULT 7

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R:Ng, W.V.; Kennedy, S.P.; Mahairas, G.G.; Berquist, B.; Pan, M.; Shukla, H.D.; Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, D.W.; Maddocks, D. Jung, K.H.; Alam, M.; Freitas, T.

Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000

A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.; Ebhardt, H.; Lowe A;Title: Genome sequence of Halobacterium species NRC-1.

A;Reference number: A84160; MUID:20504483; PMID:11016950

A;Accession: D84181
                                                                                                                               Nucleic Acids Res. 28, 4317-4331, 2000

Nucleic Acids Res. 28, 4317-4331, 2000

A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans

A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans
                                                                                                                                                                                                                                                       hypothetical protein BH0384 [imported] - Bacillus halodurans (strain C-125) C;Species: Bacillus halodurans C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 09-Jul-2004 C;Accession: H83697
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A;Title: Identification of the operon for the sorbitol (glucitol) pl A;Reference number: Z25439; MUID:20107103; PMID:10639465
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A; Residues: 1-149 <STO>
A; Cross-references: UNI
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respectes: Halobacterium sp. NRC-
                  A; Experimental source: C; Genetics:
                                    A;Cross-references: UNIPROT:Q9KFU2; UNIPARC:UPI00000C386B; GB:AP001508; GB:BA000004; NIDA;Experimental source: strain C-125
                                                                             A; Molecule type: DNA
A; Residues: 1-257 <STO>
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;Date: 18-Aug-2000 #sequence_revision 18-Aug-2000 #text_change 09-Jul-2004;Accession: T51719
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                                                                                                                          ;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;Residues: 1-187 <BOY>
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Pred. No. 41;
1; Mismatches
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Pred. No.
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G.; Jabl
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RESULT 12 AD2899

conserved

hypothetical

protein

Atu2629

[imported]

Agrobacterium

tumefaciens (strain

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A;Introns: 43/2; 92/1; 186/2
A;Note: AA TM018A10.12
C;Superfamily: Arabidopsis tl
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T01568
                                                                                                       A;Gene: AGR C_4'o,
A;Map position: circular chromosome
                                                                                                                                                                                                                        A;Reference number: A97359; WUID:21608551; PMID:11743194
A;Accession: E97674
                                                                                                                                                                                                                                                                                                                         hypothetical protein AGR_C_4767 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens C;Species: Agrobacterium tumefaciens O;Species: Acsep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004 C;Accession: E97674
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E97674
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A; Residues: 1-365 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  submitted to the EMBL Data Library, July 1997
A;Description: The sequence of A. thaliana TM018A10
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Best Local S
Matches 5
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195
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WMLSSF 200
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Pred. No.
                                                                           Score 31;
Pred. No.
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                                                                                                                                                                                                                                                                                            S.; Miller, N.; Blanchard, M.; Qurollo, M.; Doughty, D.; Scott, C.; Lappas, C.;
                                                                                                                                                                      UNIPARC: UPI00000D28EF; GB: AE007869;
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Mismatches
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Markelz,
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rkelz, B.;
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C;Species: -y--2002 #Bequence_C;Date: 11-Jan-2002 #Bequence_C;Date: 11-Jan-2002 #Bequence_C;Accession: AD2899
C;Accession: AD2899
R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Kutyavin, T. erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T. erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Karp, P.; Romero, P.; Zhang, S.
crience 294, 2317-2323, 2001
**Crience 294, 2317-2323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;de Boer, H.A., 1987
Gene 60, 93-102, 1987
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Nucleic Acids Res. 19, 599-603, 1991
A;Title: Genomic organization of lignin peroxidase genes
A;Reference number: S13563; MUID:91187681; PMID:2011531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Zhang, Y.Z.; Reddy, C.A.; Rasooly, A. Gene 97, 191-198, 1991. Respectively. Representation of several lignin peroxidase (LIP)-encoding genes: sequence analysis A;Title: Cloning of several lignin peroxidase (LIP)-encoding genes: sequence analysis A;Reference number: JN0117; MUID:91153647; PMID:1999283
                                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI000004CB03;
R;Dass, S.B.; Reddy, C.A.
FEMS Microbiol. Lett. 69, 221-224, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI00004CB03; GB:X55343; EMBIA;Note: the nucleotide sequence was submitted to the EMBL R;de Boer, H.A.; Zhang, Y.Z.; Collins, C.; Reddy, C.A.
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                                                                                                                                                    A; Reference number: A60995
A; Accession: E60995
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A; Residues: 1-371 <GAS>
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A; Residues: 1-368 <K
A;Cross-references: UNIPARC:UPI00001721E6
A;Experimental source: ATCC 24725
                                                                            A; Molecule type: protein A; Residues: 28-37 < DAS>
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                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
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;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004
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Experimental source: strain C58 (I
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                                                                                                                                                                                                                                                                                                                                                                                                  1-371 <DEB>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WMLSAF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 5; Conserv
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Pred. No. 79;
1; Mismatches
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EMBL Data Library, August 1990
                                                                                                                                                                                                                                         produced
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C;Comment: Lignin peroxidases play a key role in the initial depolymerization (;Genetics: A;Gene: GLG5; LIP6
A;Introns: 21/3; 71/2; 90/1; 161/1; 175/1; 201/2; 243/2; 343/3; 365/2
C;Superfamily: peroxidase
C;Keywords: extracellular protein; glycoprotein; heme; oxidoreductase
F;11-18/Domain: signal sequence #status predicted <SIG>
F;19-27/Domain: propeptide #status predicted <SIG>
F;19-27/Domain: propeptide #status predicted <PRO>
F;28-371/Product: lignin peroxidase H10 #status experimental <MAT>
F;283/Binding site: carbohydrate (Asn) (covalent) #status predicted
                                                                               C;Keywords: extracellular protein; glycoprotein; heme; oxide F;1-21/Domain: signal sequence #status predicted <SIG> F;22-28/Domain: propeptide #status predicted <PRO> F;29-371/Product: lignin peroxidase #status predicted <MAT> F;285/Binding site: carbohydrate (Asn) (covalent) #status p
                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Dass, S.B.; Reddy, C.A. FEMS Microbiol. Lett. 69, A;Title: Characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 1-371 <RT2>
A;Cross-references: UNIPARC:UPI000012E681; GB:M74229; NID:g169271;
A;Experimental source: strain OGC101
A;Accession: PS0247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene 107, 119-126, 1991
A;Title: Lignin peroxidase from the basidiomycete Phanerochaete chrysosporium
**Pafarence number: JH0491; MUID:92077421; PMID:1743510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gene 118, 73-80, 1992
A;Title: Characterization of a highly expressed lignin
A;Reference number: JC1268; MUID:92380494; PMID:1511887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Phanerochaete chrysosporium
C;Date: 30-Sep-1993 #sequence revision 30-Sep-1993
C;Accession: JC1268; JH0491; PS0247; C60995
R;Ritch Jr., T.G.; Gold, M.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
JC1268
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                                                                                                                                                                                                                                                                                                A;Cross-references: UNIPARC:UPI0000175215
A;Experimental source: ATCC 24725
                                                                                                                                                                                                                                                                                                                                           A; Molecule type:
A; Residues: 29-38
                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: A60995
A; Accession: C60995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-371 < RIT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Accession: JH0491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R; Ritch Jr.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Experimental source: strain OGC101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Cross-references: UNIPROT: P49012; UNIPARC: UPI000012E681;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                       ;Gene: LG2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                          Superfamily:
                                                                                                                                                                                                                                                         Genetics
                                                                                                                                                                                                                                                                              Comment: Lignin peroxidase
                                        Query Match
Local Similarity
mes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           r., T.G.; Nipper, V.J.; Akileswaran, L.;
119-126, 1991
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                                                                                                                                                                                                            21/1; 72/2; 91/2; 163/1; 177/1; 203/2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-48 <RIT1>
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                  81.6%;
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Pred. No.
Score 31; DB 2;
Pred. No. 80;
0; Mismatches
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                                           Length 371;
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lignin peroxidase (EC 1.11.1.-) LIPA precursor - basidiomycete (Phanerochae N;Alternate names: ligninase LIPA C.Species: Phanerochaete chrysosporium C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 12-Jul-2004 C;Accession: S13563
                                                                                                                                                                                                      C;Superfamily: peroxidase
C;Keywords: extracellular protein; glycoprotein; heme; oxidoreductase
C;Keywords: extracellular protein; glycoprotein; heme; oxidoreductase
C;Keywords: extracellular protein; glycoprotein; heme; oxidoreductase
C;LPA #status predicted <MAT>
F;29-372/Product: lignin peroxidase LIPA #status predicted <MAT>
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Nucleic Acids Res. 18, 7173, 1990
A; Title: Nucleotide sequence of a new lignin peroxidase A; Reference number: S13723; MUID:91088334; PMID:2129560
A; Accession: S13723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Phanerochaete chrysosporium
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 12-Jul-2004
C;Accession: S13723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Keywords: extracellular protein; glycoprotein; heme; oxidoreductase F;1-28/Domain: signal sequence #status predicted <SIG> F;1-28/Domain: signal sequence #status predicted <SIG> F;29-372/Product: lignin peroxidase GIG3 #status predicted <MAT> F;285/Binding site: carbohydrate (Asn) (covalent) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lignin peroxidase (EC 1.11.1.-) GLG3 precursor - basidiomycete (Phanerochaete chrysospor N;Alternate names: ligninase LIPA C;Species: Phanerochaete chrysosporium
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                                                                                                                                                                                                                                                                                            A;Introns: 21/1; 72/2; 91/2; 163/1; 177/1; 203/2; 344/3; 366/2 C;Superfamily: peroxidase
                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-372 < GAS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Gaskell, J.; Dieperink, E.; Cullen, D.
Nucleic Acids Res. 19, 599-603, 1991
A;Title: Genomic organization of lignin peroxidase genes
A;Reference number: S13563; MUID:91187681; PMID:2011531
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A; Residues: 1-372 <NAI>
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Local Similarity 85.7%;
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                                                                                                                                                     Score 31;
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                                                                                                           Mismatches
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lignin peroxidase (EC 1.11.1.-) GLG6 precursor - C;Species: Phanerochaete chrysosporium C;Date: 28-Jun-1991 #sequence_revision 28-Jun-199 C;Accession: A36693
                                                                                      RESULT
A36693
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R;Smith, T.L.; Schalch, H.; Gaskell, J.; Covert, S., Current, J.; Nucleic Acids Res. 16, 1219, 1988
A;Title: Nucleotide sequence of a ligninase gene from Phanerochaete A;Reference number: S01028; MUID:88144011; PMID:3344218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N;Alternate names: lignin peroxidase H8; ligninase H8 C;Species: Phanerochaete chrysosporium C;Date: 28-Feb-1990 #sequence_revision 28-Feb-1990 #text_change C;Accession: S01028
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C;Species: Phanerochaete chrysosporium
C;Date: 31-Mar-1993 #sequence_revision 31-Mar-1993
C;Accession: S13564
                                                                                                                                                                            A/Gene: LPOA
A/Introns: 21/1; 72/2; 91,
C/Superfamily: peroxidase
C/Keywords: extracellular
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A; Residues: 1-372 <SMI>
A; Cross-references: UNIPROT: P06181;
A; Note: the authors translated the C; Genetics:
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A, Title: Genomic organization of lignin peroxidase genes of A, Reference number: S13563; MUID:91187681; PMID:2011531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Keywords: extracellular protein; glycoprotein; heme; oxidoredu F;1-88/Domain: signal sequence #status predicted <SIG> F;29-372/Product: lignin peroxidase LIDB #status predicted <MAT> F;285/Binding site: carbohydrate (Asn) (covalent) #status predic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Genetics:
A;Gene: LIPB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-372 < GAS>
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N,Alternate names: lignin peroxidase H8; ligninase H8
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28-Jun-1991

#text_change

basidiomycete (Phanerochaete

chrysospor

R;Naidu,

P.S.;

Zhang, Y.Z.; Reddy, C.A.

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lignin peroxidase (EC 1.11.1.-) 48 precursor - basidiomycete (Phanerochaete chrysosporium C;Species: Phanerochaete chrysosporium C;Date: 11-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 12-Jul-2004 C;Accession: A43638 R;Andrawis, A.; Pease, E.A.; Kuan, I.; Holzbaur, E.; Tien, M. Biochem. Biophys. Res. Commun. 162, 673-680, 1989 A;Title: Characterization of two lignin peroxidase clones from Phanerochaete chrysospori A;Reference number: A43638; MUID:89334875; PMID:2474293 A;Accession: A43638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Walther, I.; Kaelin, M.; F
Gene 70, 127-137, 1988
A; Title: Molecular analysis
A; Reference number: JT0402;
A; Accession: JT0402
                                                                                                  A;Cross-references: UNIPARC:UPI0000168966; GB:M27884; NID:g169283; PIDN:AAB00798.1; A;Note: the authors translated the codon TGC for residue 41 as Ser, and TCG for resic;Superfamily: peroxidase C;Kuperfamily: peroxidase
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C;Superfamily: peroxidase
C;Keywords: extracellular
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A;Residues: 1-372 <WAL>
A;Cross-references: UNIPROT:P06181; UNIPARC:UPI000012E685
C;Genetics:
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A;Title: Characterization of a new lignin peroxidase gene (GLG6) from Phanerochaete
A;Reference number: A36693; MUID:91097609; PMID:2268360
A;Accession: A36693
                                                                                                                                                                                 A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-372 <AND>
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C;Species: Phanerochaete chrysosporium
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Similarity 85.7%;
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                                                              DB 2;
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lignin peroxidase (EC 1.11.1.-) H8 precursor - NyAlternate names: ligninase H8 C;Species: Phanerochaete chrysosporium C;Date: 27-Feb-1990 #sequence_revision 31-Dec-1
                                                                               RESULT
A32322
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A; Gene: liP
C; Function:
A; Description: 1
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R;Schneider, H.; Barth, W.; Boehme, H.J.

Biol. Chem. Hoppe-Seyler 377, 399-402, 1996

A;Title: Cloning and characterization of another lignin
A;Reference number: S69246; MUID:96437179; PMID:8839987

A;Accession: S69246
                                                                                                                                           A;Cross-references: UNIPROT:Q01775; UNIPARC:UPI0000175213 C;Superfamily: peroxidase C;Keywords: extracellular protein; glycoprotein; heme; ox
                                                                                                                                                                                                                                                                              R;Andrawis, A.; Pease, E.A.; Kuan, I.; Holzbaur, E.; Tien, M. Biochem. Biophys. Res. Commun. 162, 673-680, 1989
A;Title: Characterization of two lignin peroxidase clones from A;Reference number: A43638; MUID:89334875; PMID:2474293
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A; Residues: 1-372 < AND>
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A;Experimental source: strain BKM-F 1767
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                                                                                                                                                                                                                                                                                                                                                            ;Species: Phanerochaete chrysosporium;Date: 11-Feb-1993 #sequence_revision;Accession: B43638
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WMLSAHS
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31-Dec-1993 #text_change 12-Jul-2004

basidiomycete (Phanerochaete

chrysosporium

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RESULT 26
B32322
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R;Dass, S.B.; Reddy, C.A.
FEMS Microbiol. Lett. 69, 221-224, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI0000175221; GB:M27400
A;Note: the authors translated the codon CAG for residue 11
R;Doyle, W.A.; Smith, A.T.
Biochem. J. 315, 15-19, 1996
A;Title: Expression of lignin peroxidase H8 in Escherichia
A;Reference number: 866556; MUID:96207556; PMID:8670100
A;Accession: 866556
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A;Title: Molecular cloning and sequences of lignin peroxidase genes of Phanerochaete chr
A;Reference number: A32322; MUID:89343994; PMID:2761543
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R;Schalch, H.; Gaskell, J.; Smith, T.L.; Cullen,
Mol. Cell. Biol. 9, 2743-2747, 1989
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A;Title: Multiple ligninase-related genes from Phanerochaete chrysosporium
A:Reference number: JS0065; MUID:89211962; PMID:2907500
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A; Residues: 1-372 <S
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A;Residues: 1-372 <BRO>
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R;Brown, A.; Sims, P.F.G.; Raeder, U.; Broda, P.
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A;Experimental source: ATCC 24725
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A;Residues: 29-38 <DAS>
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A;Accession: D60995
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A;Residues: 22-30 <DOY>
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                                                                                                                                                                                                                                                           ;Superfamily: peroxidase;Reywords: extracellular protein; glycoprotein; heme; oxidoreductase;1-28/Domain: signal sequence #status predicted <SIG>
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Similarity 85.7%;
6; Conservation
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C;Accession: B32322
R;Schalch, H.; Gaskell, J.; Smith, T.L.; Cullen, D.
Mol. Cell. Biol. 9, 2743-2747, 1989
A;Title: Molecular cloning and sequences of lignin peroxidase
A;Reference number: A32322; MUID:89343994; PMID:2761543
A;Accession: B32322
A;Accession: B32322
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C;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 12-Jul-2004
C;Accession: JH0156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lignin peroxidase (EC 1.11.1.-) 0282 precursor - basidiomycete (Phanerochaete C;Species: Phanerochaete chrysosporium C;Date: 27-Feb-1990 #sequence_revision 27-Feb-1990 #text_change 12-Jul-2004
                                                                                                                                                                                                                                                                     R;Tien, M.; Tu, C.P.D.
Nature 326, 520-523, 1987
A;Title: Cloning and sequencing of a cDNA in the control of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Phanerochaete chrysosporium C;Date: 16-Aug-1988 #sequence_revision C;Accession: A27817
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A;Title: Characterization of lignin peroxidase-encoding
A;Reference number: JH0156; MUID:90323600; PMID:2373364
F;1-28/Domain: signal sequence #status predicted <SIG>F;29-373/Product: lignin peroxidase #status predicted
                                                                         C; Superfamily: peroxidase
C; Reywords: extracellular protein; glycoprotein; heme; oxidoreductase
                                                                                                                                                       A;Cross-references: UNIPROT:P06181; UNIPARC:UPI0000168965; GB:Y00262; NID:g3149;
                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-373 <TIE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Genetics:
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A;Residues: 1-372 <HUO>
A;Cross-references: UNIPROT:P31837; UNIPARC:UPI0000168967; GB:M37701; GB:M22720; NID:g169
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RESULT 31
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A;Title: Genome Sequence and Comparative Analysis of the A;Reference number: A96900; MUID:21359325; PMID:21359325
A;Accession: D96955
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A; Residues: 1-416 < KUR>
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transcription regulator AraC/XylS family VC2080 [imported] - Vibrio cholerae C;Species: Vibrio cholerae C;Species: Vibrio cholerae C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004 C;Accession: C82120
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A;Experimental source: cosmid contig 68B2; strain 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Schulte, U.; Aign, V.; Hoheisel, J.; Brandt, P.; Fartmann, submitted to the Protein Sequence Database, April 2000
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C;Date: 05-May-2000 #sequence_revision 05-May-2000 #text_change
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A;Residues: 1-541 <SCH>
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A;Accession: T48836
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:Accession: D96955
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nilarity 85.7%;
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Pred. No. 1.1e+02;
1; Mismatches 0;
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                                                                                                                                                                                                                                                Length 541;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B.; Holland, R.;
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A; Introns: (C; Superfami)
C; Keywords:
hypothetical protein SPAC17C9.10 - fission yeast (SC;Species: Schizosaccharomyces pombe C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 C;Accession: T11590; T52026 R;Barrell, B.G.; Rajandream, M.A.; Walsh, S.V. submitted to the EMBL Data Library, April 1996 A;Reference number: Z17295
                                                                                                                                               RESULT 33
T11590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J.; Chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, P. R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MUID:20406833; PMID:10952301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Residues: 1-1121 <HEI>
A;Cross-references: UNIPROT:Q9KQCO; A;Experimental source: serogroup O1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Accession: B53189; S41020
R; Hengartner, M.O.; Horvitz, H.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytochrome b560 homolog precursor - Caenorhabditis elegans N; Alternate names: hypothetical protein T07C4.7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cell 76, 665-676, 1994
A;Title: Caenorhabditis elegans cell survival gene ced-9
A;Reference number: A53189; MUID:94170367; PMID:7907274
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A;Residues: 1-182 <B
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A; Residues: 1-182 < HEN>
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Best Local (
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Keywords: membrane protein; mitochondrion
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Cross-references: UNIPARC:UPI0000126C6F; EMBL:Z29443; NID:g1067051; PID:g443834
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Pred. No.
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Pred. No.
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09-Jul-2004

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R:Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H. ansen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 816-820, 2000
A.; Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, F. Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A.; Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A.; Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A.; Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                              RESULT 35
G64141
probable tetraacyldisaccharide 4'-kinase (EC 2.7.1.130) HI0059 [similarity] C;Species: Haemophilus influenzae C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 09-Jul-2004 C;Accession: G64141 R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.;
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C; Superfamily: P
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A; Residues: 1-271 < KYU>
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A;Experimental source: strain 972h(-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references:
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A; Residues: 1-305 <ST
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A;Experimental source: strain ED665h(-)
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Best Local S
Matches 5
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;Species: Arabidopsis thaliana (mouse-ear cress)
;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 05-Oct-2004
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71.4%;
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71.4%;
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Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                        Length 305
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                                                                                       Haemophilu
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war, K.;
                                                                                                                                                                                                                                                   R;Goodner, B.; ....R;Goodner, B.; Wollam,
                                                               A;Status: preliminary
A;Molecule type: DNA
A;Mesiduse: 1-348 «KUR»
A;Cross-references: UNIPROT:Q8UHI5; UNIPARC:UPI000012E91B;
C;Genetics:
                                                                                                                                                                                                                                              R;Goodner, B.; Hinkle, G.; Gattung, A.; Liu, F.; Wollam, C.; Allinger, Science 294, 2323-2328, 2001
A;Gene: AGR C 1257
A;Map position: ci
C;Superfamily: Esc
                                                                                                                                                                                                                                                                                                                                 C;Species: Agrobacterium tumefaciens
C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 09-Jul-2004
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                                                                                                                                                                                                      A;Title: Genome Sequence of the Plant Pathogen and Biotechnology A;Reference number: A97359; MUID:21608551; PMID:11743194
                                                                                                                                                                                                                                                                                                                     C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                         tetraacyldisaccharide 4'-kinase (lipid a 4'-kinase) [imported] - Agrobacterium
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B97444

S.; Miller, N.; Blanchard, M.; Qurollo, M.; Doughty, D.; Scott, C.; Lappas, C.;

Agent

Agrobacterium

tum

B.; Goldn Markelz,

Goldman, kelz, B.;

tumefacier

: circular chromosome Escherichia coli pro

probable

tetraacyldisaccharide

4'-kinase

ycaH

GB:AE007869;

PIDN: AAK86507.1;

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C;Accession:
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Science 269, Control of the Contr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;She, Q.; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Char Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               glycosyltransferase [imported] - Sulfolobus solfataricus
C;Species: Sulfolobus solfataricus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Residues: 1-341 < KUR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Accession: C90322
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Best Local
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WLLSLFS
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71.4%;
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Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
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                                                                                                                                                                                                                                               Length 341;
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R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel; Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm, A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.;
                                                                                                                                                                                                                                                                                                                                                          R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J. chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, I, R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MUID:20406833; PMID:10952301
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A;Experimental source: serogroup O1; strain N16961; biotype El Tor
                                                                                                                                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-374 < HEI>
                                                                                                                                                                                                                                                                                                                    A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                        A;Reference number: A82035;
A;Accession: E82168
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C;Superfamily: Escherichia coli probable tetraacyldisaccharide 4'-kinase
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A;Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Species: Vibrio cholerae
;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004
                                                                                                                                                                                      Genetics:
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                                                                                                       78.9%;
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71.4%;
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71.4%;
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Pred. No. 1.36
1; Mismatches
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Pred. No. 1.2e+02;
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Pred. No.
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1.3e+02;
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C;Accession: S65138; G48394
R;Aoki, N.; Kishi, M.; Taniguchi, Y.; Adachi, T.; Nakamura, R.; Matsuda, T. Biochim. Biophys. Acta 1245, 385-391, 1995
A;Title: Molecular cloning of glycoprotein antigens MGP57/53 recognized by monoclonal A;Reference number: S65138; MUID:96125736; PMID:8541316
A;Accession: S65138
A;Status: preliminary; not compared with conceptual translation
                                                                                                                                   A;Note: sequence extracted from NCBI backbone C;Superfamily: milk fat globule protein; disco C;Keywords: glycoprotein F;1-32/Domain: BGF homology (fragment) <EG1>F;40-79/Domain: BGF homology <EG2>F;82-239/Domain: discoidin I amino-terminal ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI0000177B00
R;Matcher, I.H.; Banghart, L.R.; Lane, W.S.
Biochem, Mol. Biol. Int. 29, 545-554, 1993
A;Title: The major fat-globule membrane proteins, bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      glycoprotein antigen MGP57/53, mammary gland - bovine (fragment)
N,Alternate names: glycoprotein component 16/major fat-globule membrane protein/MFG-E8
C;Species: Bos primigenius taurus (cattle)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 07-Aug-1998
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S65138
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A;Map position: 2
C;Superfamily: tyrosine-specific transport protein
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                                                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: protein
A;Molecule type: protein
A;Residues: 207-220 <MAT>
A;Cross-references: UNIPARC:UPI000014C31B
                                                                                                                                                                                                                                                                                                 A; Experimental source: milk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               II-like sequences.
A; Reference number: A48394; MUID:93250576; PMID:8485470
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                                                                                                         ;82-239/Domain: discoidin I amino-terminal;243-401/Domain: discoidin I amino-terminal
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                       78.9%;
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I amino-terminal homology <DN2</pre>
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                       Score 30; DB 2; 1
Pred. No. 1.4e+02;
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Pred. No. 1.
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H.; Dragoi, I.; Sellers,
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A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; Lelaure, A;Authors: Kahn, M.L.; Kalman, B;Authors: Kahn, M.L.; Kal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Abola, A.P.; Barloy-Hubler, J. Kalman, S.; Keating, D.H.; Palm, C.; Peck, M.C.; Surzycki, R.; Wells, D.H.; Proc. Natl. Acad. Sci. U.S.A. 98, 9883-9888, 2001
A;Tile: Nucleotide sequence and predicted functions of the entire Sinorhizobiu A;Reference number: A95262; MUID:21396509; PMID:11481432
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C; Superfamily:
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A; Residues: 1-412 < KUR>
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P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K.; Yeh, K
The composite genome of the legume symbiont Sinorhizobium meliloti.
nce number: A96039; MUID:21368234; PMID:11474104
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Biochem. Mol. Biol. Int. 29, 5. A; Title: The major fat-globule
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Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant
A;Reference number: ABO502; MUID:21534947; PMID:11677608
A;A;Ccession: ACO546
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eur. J. Biochem. 240, 628-636, 1996
A;Title: Characterization of glycoprotein PAS-6/7 from membranes of bovine milk fat A;Reference number: $74211; MUID:97008954; PMID:8856064
A;Accession: $7421
A;Molecule type: mRNA
A;Residues: 1-427 <HVA>
                                                                                                                                                                                                       A;Cross-references: UNIPARC:UPI000017780B
R;Aoki, N.; Kishi, M.; Taniguchi, Y.; Adachi, T.; Nakamura, R.; Matsuda, Biochim. Biophys. Acta 1245, 385-391, 1995
A;Title: Molecular cloning of glycoprotein antigens MGP57/53 recognized A;Reference number: S65138; MUID:96125736; PMID:8541316
                                                                                                                                                                                                                                                                                                                                                                                                                  R;Kim, D.H.; Kanno, C.; Mizokami, Y.
Biochim. Biophys. Acta 1122, 203-211, 1992
A;Title: Purification and characterization of major glycoproteins, A;Reference number: S23926; MUID:92353107; PMID:1643094
A;Accession: S24181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 19-85;96-110;140-165;174-216;221-232;248-277;285-293;309-337;339-420;425-427
A;Cross-references: UNIPARC:UPI0000177B01; UNIPARC:UPI0000177B02; UNIPARC:UPI0000177B03;
B08; UNIPARC:UPI0000177B09; UNIPARC:UPI0000177B0A
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C;Date: 04-Dec-1997 #sequence revision 12-Dec-1997 #text_change
C;Accession: S74211, S78114, S24181, S65138; G48394
R;Hvarregaard, J.; Andersen, M.H.; Berglund, L.; Rasmussen, J.T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N;Alternate names: glycoprotein component 16/major fat-globule membrane protein/MFG-E8
A;Cross-references: UNIPARC:UPI0000177B00 R;Mather, I.H.; Banghart, L.R.; Lane, W.S. Biochem. Mol. Biol. Int. 29, 545-554, 1993
                                                                                         A; Molecule type: mRNA
A; Residues: 27-427 < AOK>
                                                                                                                                                A; Status: preliminary; not
                                                                                                                                                                               A; Accession: S65138
                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: protein A; Residues: 383-394 < KIM>
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A; Residues: 1-417 < PAR >
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83.3%;
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Pred. No. 1.4e+02;
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membrane proteins,

bovine components 15/16

and

guinea-pig

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C;Accession: A75444
C;Accession: A75444
R;Mhite, O.; Eisen, J.A.; Heidelberg, J.F.; Hickey, E.K.; Peterson, M.; Shen, M.; Vamathevan, J.J.; Lam, P.; McDonald, L.; Utterback, S.; Smith, H.O.; Venter, J.C.; Fraser, C.M.
Science 286, 1571-1577, 1999
A;Title: Genome sequence of the radioresistant bacterium Deinococcus A;Tit
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A;Accession:
A;Status: pre
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F;66-105/Domain: EGF homology <EG1>
F;108-265/Domain: EGF homology <EG2>
F;108-265/Domain: discoidin I amino-terminal homology <DN1>
F;269-427/Domain: discoidin I amino-terminal homology <DN2>
F;24-35,29-47,49-58,66-77,71-94,96-105/Disulfide bonds: #status predicted F;24-35,29-47,49-58,66-77,71-94,96-105/Disulfide bonds: #status experimental F;27/Binding site: carbohydrate (Ser) (covalent) #status experimental F;34/Binding site: carbohydrate (Thr) (covalent) #status experimental F;59,227/Binding site: carbohydrate (Asn) (covalent) #status experimental F;59,227/Binding site: carbohydrate (Asn) (covalent) #status experimental
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C;Superfamily: milk fat globule protein; discoidin I amino-terminal homology;
C;Keywords: blocked amino end; disulfide bond; glycoprotein; milk
F;1-18/Domain: signal sequence #status predicted <SIG>
F;19-427/Product: PAS-6/7 protein #status experimental <MAT>
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C;Species: Deinococcus radiodurans
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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A;Molecule type: prottein
A;Residues: 233-246 <MAT>
A;Cross-references: UNIPARC:UPI000014C31B
                                                                                                                                                          conserved hypothetical protein MYPU_2620 [imported] - Mycoplasma pulmonis (strain UAB C;Species: Mycoplasma pulmonis C;Species: Mycoplasma pulmonis C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 09-Jul-2004 C;Accession: F90544
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C; Superfamily:
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A; Residues: 1-454 < WHI>
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R;Chambaud, I.; Heilig, R.; Ferris, S.; Barbe, V.; Samson, D.; Galisson, F.; Moszer, Nucleic Acids Res. 29, 2145-2153, 2001
Nucleic Acids Res. 29, 2145-2153, 2001
A;Title: The complete genome sequence of the murine respiratory pathogen Mycoplasma A;Reference number: A99512; MUID:21267165; PMID:11353084
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Pred. No. 1.5e+02;
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Pred. No. 1.6e+02;
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                                                                                                                    D.; Galisson, F.; Moszer,
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T.; Zalewski,
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submitted to the EMBL Data Library, May 1998 A;Reference number: Z21939 A;Accession: TAGECO
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R;Blanc, V.; Salah-Bey, K.; Folcher, M.; Thompson, C.J.

Mol. Microbiol. 17, 989-999, 1995

A;Title: Molecular characterization and transcriptional analysis of a multidrug resistance number: S70171; MUID:96123441; PMID:8596448

A;Accession: S70171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Gene: MYPU 2620
A;Genetic code: SGC3
C;Superfamily: hypothetical protein MG294
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A;Residues: 1-467 <KUR>
A;Cross-references: UNIPROT:Q98QU9; UNIPARC:UPI00000C8046; GB:AL445566; PID:g14089675;
A;Experimental source: strain UAB CTIP
                                                                                                                                                                                                                                                                                                                                                              hypothetical protein SPBC649.05 - fission yeast (Schizosaccharomyces pombe) C;Species: Schizosaccharomyces pombe C;Date: 03-Dec1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004 C;Accession: T40598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-501 <BLA>
A;Cross-references: UNIPROT:Q54806; UNIPARC:UPI00000B6054; EMBL:X84072; NID:g872305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-467 < K
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                                                                                                                                                                                       A;Cross-references: UNIPROT:059755; UNIPARC:UPI0000128686; EMBL:AL023587; PIDN:CAA19047.1
                                                                                                                                                                                                                                            A; Molecule type:
                                                                                                                                                                                                                                                              A;Status: preliminary; translated from GB/EMBL/DDB
                                                                                                                                                                                                                                                                                  A; Accession: T40598
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Pred. No. 1.9e+02;
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Pred. No. 1.7e+02;
3; Mismatches 0
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Pred. No. 1.6e+02;
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WVLKAFS

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RESULT 50
A332280
glutamate receptor GluR-II precursor - fruit fly (Drosophila melanogaster)
C;Species: Drosophila melanogaster
C;Date: 28-Feb-1992 #sequence_revision 10-Apr-1992 #text_change 31-Dec-2004
C;Accession: A39280
R;Schuster, C.M.; Ultsch, A.; Schloss, P.; Cox, J.A.; Schmitt, B.; Betz, H.
R;Schuster, C.M.; Ultsch, A.; Schloss, P.; Cox, J.A.; Schmitt, B.; Betz, H.
R;Schuster, C.M.; Ultsch, A.; Schloss, P.; Cox, J.A.; Schmitt, B.; Betz, H.
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R;Schuster, C.M.; Ultsch, A.; Schloss, P.; Cox, J.A.; Schmitt, B.; Betz, H.
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Title: Perfect score: Run on: OM protein - protein search, using sw model August 29, 2006, 05:47:27; Search time 56 Seconds (without alignments)
115.627 Million cell updates/sec US-10-541-343-3 38 Copyright GenCore version 5.1.9 (c) 1993 - 2006 Biocceleration Ltd.

Searched: Scoring table: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 WMLSAFS 7

2849598 seqs, 925015592 residues

Total number of hits satisfying chosen parameters:

2849598

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 500 summaries Minimum DB Maximum DB seq length: 0
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UniProt_7.2:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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InterPro; IPR00337; GPCR Mgr.
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Pfam; PF00003; 7tm 3; 1.
Pfam; PF007562; NCD3G; 1.
PFAM; PF07562; NCD3G; 1.
PROSITE; PS00980; G PROTEIN RECEP F3 2; 1.
PROSITE; PS00981; G PROTEIN RECEP F3 3; 1.
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RECEPTOR: Transmembrane.
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Takami H., Takaki Y., Chee G.-J., Nishi
Matsui S., Uchiyama I.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tsutsumi S., Aburatani H., Asai K., Akiyama Y. Submitted (JUL-2001) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suwa M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2002, sequence version 1.
07-FEB-2006, entry version 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QBNOZO_HUMAN
                                                                                                                                                                                                                                                                Bacteria; Firmicutes; Bacillales;
                                                                                                                                                                                                                                                                                       Geobacillus kaustophilus
                                                                                                                                                                                                                                                                                                                      OrderedLocusNames=GK0704;
                                                                                                                                                                                                                                                                                                                                          Maltose/maltodextrin transport system (Subsrate-binding protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Suwa M., Sato
Tsutsumi S., I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seven transmembrane helix receptor.
                                                "Thermoadaptation trait revealed by the thermophilic Geobacillus kaustophilus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human)
                         Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GEOKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AB065450; BAC05713.1; -; Genomic_DNA.
AB065566; BAC05802.1; -; Genomic_DNA.
AB065486; BAC05739.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 WMLSAFS 7
                         Acids Res. 32:6292-6303(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WMLSSFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENSG00000198822; Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T., Okouchi I., Arita M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA; 39300 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                [LARGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92.1%;
85.7%;
                                                                                                                                                                                             SCALE GENOMIC DNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 35; DB
Pred. No. 75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B114CF38477FF624 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                     Bacillaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     424 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Futami K.,
                                                                     genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GABA-B-like recepto.
                                                                                                                        Shimamura S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 347;
                                                                       sequence
                                                                                                                                                                                                                                                                  Geobacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matsumoto
                                                                            얁
                                                                                                                        Suzuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IEA
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QUESTLATION OF THE COLOR OF THE
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                                                                                                    R GO; GO:0005739; C:mitochondrion; IEA.

R GO; GO:0008137; F:NADH dehydrogenase (ubiquinone) activity; IEA

R GO; GO:0016491; F:oxidoreductase activity; IEA.

R GO; GO:0016491; F:oxidoreductase activity; IEA.

R GO; GO:00042773; P:ATP synthesis coupled electron transport; IEA

R GO; GO:0006120; P:mitochondrial electron transport, NADH to u.

R InterPro; IPR003918; NADHub oxred4.

R InterPro; IPR001750; Oxidored q1.

R InterPro; IPR001750; Oxidored q5.

R InterPro; IPR000260; Oxidored q5.

R PANNTHER; PTHR11460; NADHub oxred4; 1.

R Pfam; PF00361; Oxidored q5.

R Pfam; PF01059; Oxidored q5.

R Pfam; PF01059; Oxidored q5.

R PRINTS; PR01437; NUOXDRDTASE4.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Dutheria; Lucrdata; Craniata; Vertebrata; Eut
Hystricognathi; Thryonomyidae; Thryonomys.
NCBI TaxID=10169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; BA000043; BAD74989.1; -; Genomic_DNA.
GO; GO:0005363; F:maltose transporter activity; IEA.
GO; GO:0005215; F:transporter activity; IEA.
GO; GO:0015768; P:maltose transport; IEA.
GO; GO:0016768; P:transport; IEA.
GO; GO:0006810; P:transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0006810; P:transport; IEA.
InterPro; IPR006060; Maltose_bd.
InterPro; IPR006059; SBP_bac_1.
InterPro; IPR006059; SBP_domI.
Pfam; PF01547; SBP_bac_1; 1.
PRINTS; PR00181; MĀLTOSĒBP.
PROSITE; PS01037; SBP_BACTERIAL_1; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dehydrogenase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NADH-ubiquinone oxidoreductase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2001, integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9B6F3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9B6F3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Complete proteome. SEQUENCE 424 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
       TIGRFAMs; TIGR01972; NDH I_M; 1.
Mitochondrion; NAD; Oxidoreductase; Ubiquinone.
SEQUENCE 459 AA; 51966 MW; E6DD0150CC0C9F56 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AJ301644; CAC27808.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thryonomys swinderianus (Greater cane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=NADH4;
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85.7%;
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RA Adams M.D., Celniker S.E., Hilt R.A., Evans C.A., Gocayne J.D.,
RA George R.A., Lewis S.E., Richards S., Ashburner M.A., Galale R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Abril J.F., Agbayani A., An H.-J., Andrews-Ffamkoch C., Baldwin D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Baldwin D.,
RA Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Boltshakov S.,
RA Berkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Berkova D., Botchan M.R., Bouck J., Brokstein P., Brottier S.M.,
RA Gerry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.B., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Hostin D., Houston K.A., Howland T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Heimson J.A., Ketchum K.A.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Indepwan C.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Menson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb JM.,
RA Rainert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Styong A.H., Wang X.,
RA Sheng X.H., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
Welson D.R., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
Welson E.R., Myers E.W., Kubin G.M., Venter E., Wang A.H., Wang X.,
RA Sheng X.H., Zhong F., Rabin G.M., Venter J.C.;
RA Sheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu X., Smith H.O.,
RA Sheng X.H., Zhong W., Zhou X., Zhu X., Smith H.O.,
RA Sheng X.H., Wang S., Wang S., Zhu X., Smith H.O.,
RA Sheng X.H., Wang S., Wang S., Zhou X., Smith H.O.,
RA Sheng R
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Q9VLT3;
01-MAY-2000, i
                         George R.A.,
Pacleb J.M.,
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Ephydroidea; Drosophilidae;
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                                                                                                                          Celniker S.
                                                                                                                                                      MEDLINE=22426065;
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                            S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Ha
Adams M., Champe M., Dugan S.M., Frise E., Hodgson
A., Hoskins R.A., Laverty T., Mizny D.M., Nelson C.
A., Park S., Pfeiffer B.D., Richards S., Sodergren
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Endopterygota; Diptera; Brachycera; Muscomorpha;
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                                                                                                                                                   PubMed=12537568;
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                                                           E., Hodgson A.
      G.G.,
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Pfam; PF00207; A2M; 1.
Pfam; PF07678; A2M comp; 1.
Pfam; PF01835; A2M N; 1.
Pfam; PF07703; A2M N; 2; 1.
Pfam; PF0777; A2M recep; 1.
Pfam; PF0057; Ldl recept a; 1.
SMART; SM00192; LDLa; 1.
                                                                                                                                                                                                                                                                                                                                                         Stapleton M., Brokstein P., Hong L., Agbayani A., C: Champe M., Chavez C., Dorsett V., Farfan D., Frise I Gonzalez M., Guarin H., Li P., Liao G., Miranda A., Nunoo J., Pacleb J., Paragas V., Park S., Phouanena Yu C., Lewis S.E., Rubin G.M., Celniker S.; Submitted (Comm. Com.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Berkeley Drosophila Genome Project;
Celniker S., Carlson J., Wan K., Pf
Hoskins R., Stapleton M., Pacleb J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Annotation of the Drosophila melanogaster systematic review.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S. Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E. Smith C.D., Tupy J.L., Whitfield B. Bayraktaroglu L., Berman E Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A., Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q., Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDITIDE SEQUENCE.
MEDITIDE SEQUENCE.
MEDITIDE 22426069; PubMed=12537572;
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"The transposable elements of the a genomics perspective.";
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Genome Biol.
           SMART; SM00192;
PROSITE; PS50068
                                                                                                                                     InterPro; IPR002890; A2M_N.
InterPro; IPR011625; A2M_N_Z
InterPro; IPR011627; A2M_Ycpt.
InterPro; IPR002172; LDL_rcpt_
InterPro; IPR002172; LDL_rcpt_
                                                                                                                                                                                            FlyBase; FBgn0020240; Mcr.
GO; GO:0005515; F:protein binding;
InterPro; IPR011626; A2M_comp.
                                                                                                                                                                                                                                                                EMBL; AE003619; AAF52601.1; -; EMBL; AY058526; AAL13755.1; -;
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                                                                                                                                                                                                                                                                                                                                                    Submitted
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Patel S., Frise E., Wheeler D.A., Lewis S.E.,
                                                                                                                           InterPro;
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                                                                                                                           IPR001599;
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1. 3:RESEARCH0079-RESEARCH0079(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE.
                                                                                                                                                                                                                                                                                                                         by the UniProt Consortium,
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               LDLRA
                                                                                                                         LDL_rcpt_A.
MacrogloblnA2.
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.RA_2; 1.
202783 MW;
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the EMBL/GenBank/DDBJ
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5 J., Park
 B8E0602350E87F86 CRC64;
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                                                                                                                                                                                                                                                                                                          Attribution-NoDerivs License
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Svirskas 1
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Drosophila
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RESULT 6

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ID 4034A91_RHOPA
AC 034A91_AC
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DT 06-DEC-2006, e
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GN 07-FEB-2006 e
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RC STRAIN-BisB18;
RG US DOE Joint G
RA Copeland A., Ist
RT "Sequencing of
RT Palustris B1sB
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RC STRAIN-BisB18;
RG US DOE Joint G
RA Larimer F., La
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RT Submitted (NOV
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DR GG; GG:00016740
DR GG; GG:00016740
DR GG; GG:0001679
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antioxidant p
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-2005, sequence version 07-FEB-2006, entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (NOV-2005) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavin. Hammon N., Israni S., Pitluck S., Richardson P.; "Sequencing of the draft genome and assembly of Rhodopseudomonas palustris BisBl8."; Submitted (OCT-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amino acid adenylation. ORFNames=RPCDRAFT_2142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=BisB18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rhodopseudomonas palustris BisB18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Larimer F., Land M.; "Annotation of the draft genome assembly of Rhodopseudomonas"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
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                                                                                                                         BACCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOE Joint Genome Institute (JGI-ORNL);
                                                                                                                                                                                                                                                                                                                                      GO:0016874; F:ligase activity; IEA.
GO:0030170; F:pyridoxal phosphate binding;
GO:0008483; F:transaminase activity; IEA.
GO:0016740; F:transferase activity; IEA.
GO:0016633; P:fatty acid biosynthesis; IEA.
GO:0008452; P:metabolism; IEA.
GO:0008452; P:metabolism; IEA.
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                                                                                                                                                                                              1502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           preliminary data.
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 protein
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 sequence version 1.
entry version 3.
DNA-binding protein Dps
protein / ferroxidase.
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                                                                     integrated into UniProtKB/TrEMBL.
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Pred. No. 4e+02;
                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                        PRT;
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                  iron-binding
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E7 BACCZ
Q63CE7 BACCZ
Q63CE7;
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Hoffmaster A.R., Ravel J., Rasko D.K., Chapman G.D., Chut
Marston C.K., De B.K., Sacchi C.T., Fitzgerald C., Mayer
Maiden M.C.J., Priest F.G., Barker M., Jiang L., Cer R.Z.
Rilstone J., Peterson S.N., Weyant R.S., Galloway D.R., F
                                                                                                                                  Brettin T.S., Bruce D., Challacombe J.F., Gilna P., Han Hitchcock P., Jackson P., Keim P., Longmire J., Lucas S. Richardson P., Rubin E., Tice H.; "Complete genome sequence of Bacillus cereus ZK."; Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                           Name=dps; OrderedLocusNames=BCE3JL1825; Bacillus cereus (strain ZK / E3JL).
Bacteria; Firmicutes; Bacillales; Bacillascillus cereus group.
NCBI TaxID=288681;
                                                                                                                                                                                                                                                                                                                                                                                                                      25-OCT-2004, integrated into Unip
25-OCT-2004, sequence version 1.
07-FEB-2006, entry version 14.
Non-specific DNA-binding protein,
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GO; GO:0003677; F:DNA binding;
GO; GO:0008199; F:ferric iron k
GO; GO:0006879; P:iron ion home
GO; GO:0006950; P:response to g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Popovic T., Fraser C.M.;
"Identification of anthrax toxin genes in a Bacillus with an illness resembling inhalation anthrax.";
Proc. Natl. Acad. Sci. U.S.A. 101:8449-8454(2004).
-i- CAUTION: The sequence shown here is derived from the control of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus; Bacillus cereus group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=G924
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                                                                     Copyrighted
Distributed
                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRINTS; PR01346; HELNAPAPROT. ProDom; PD149803; DPS; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GO; GO:0006879; P:iron ion homeostasis; GO; GO:0006950; P:response to stress; II InterPro; IPR002177; DPS_DNA_bd.
Pfam: PF00210: Ferritin; 1.
                      EMBL; CP000001; AAU18426.1;
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Q63CE7;
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                                                                  by the UniProt Consortium, see http://www.uniprot.org/tounder the Creative Commons Attribution-NoDerivs License
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D., Challacombe J.F., Gilna
on P., Keim P., Longmire J.,
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b; Pred. No. 51;
0; Mismatches
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                        Genomic_DNA
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)kinaka R.
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ID Q6HJVJ BACHK
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ID Q6HJVJ BACH
ID Q6HJVJ BACH
ID Q6HJVJ BACH
ID 19-JUL-2
DT 19-JUL-2
DT 19-JUL-2
DT NON-Spec
GN Name-dps
OC Bacteria
OC Bactillus
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OC Bacteria
OC Bactillus
OC STRAIN-9
RA Brettin
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RA Hitchcon
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CC STRAIN-9
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DR RHI; AE
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GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0004322; F:ferroxidase activity; IEA.
GO; GO:0016491; F:oxidoreductase activity; IEA.
GO; GO:0016491; F:oxidoreductase activity; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006950; P:response to stress; IEA.
InterPro; IPR00217; DPS_DNA_bd.
InterPro; IPR008331; Ferritin_Dps.
InterPro; IPR018347; Ferritin_rel.
Pfam; PF00210; Ferritin; II.
                                                                                                                                                                      Pfam; PF00210; Ferritin; 1.
Pfam; PF00210; Ferritin; 1.
PR.NTS; PR01346; HELNAPAPROT.
ProDom; PD149803; DPS; 1.
PROSITE; PS00818; DPS; 1; 1.
PROSITE; PS00818; DPS; 1; 1.
                                                                                                                                                                                                                                                                                                                                                                                  EMBL; AE017355; AAT63428.1; -; Genomic_DNA.
SMR; O6HJV3; 2-147.
GO; GO:0003677; F:DNA binding; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0008122; F:ferroxidase activity; IEA.
GO; GO:0016491; F:oxidoreductase activity; IGO; GO:00166979; F:iron ion homeostasis; IEA.
GO; GO:0006879; P:iresponse to stress; IEA.
                                                                                                                                                      Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brettin T.S., Bruce D., Challacombe J.F., Gilna P., Han C., I Hitchcock P., Jackson P., Keim P., Longmire J., Lucas S., Ok: Richardson P., Rubin E., Tice H.; "Complete genome sequence of Bacillus thuringiensis 97-27."; Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
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PRODOM; PD149803, DPS; 1.
PROSITE; PS00818; DPS; 1.
Complete proteome; DNA-binding;
SEQUENCE 147 AA; 16707 MW; 2
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Bacillus thuringiensis subsp. konkukian.
Bacteria; Firmicutes; Bacillales; Bacilla
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19-UUL-2004, sequence version 1.
07-FEB-2006, entry version 14.
Non-specific DNA-binding protein Dps
                                                                                                                                                                                                                                                                                                      InterPro; IPR002177; DPS_DNA_bd.
InterPro; IPR008331; Ferritin_Dps.
InterPro; IPR012347; Ferritin_rel.
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GO:0008199; F:ferric iron binding; IEA.
GO:0008122; F:ferroxidase activity; IEA.
GO:0016491; F:oxidoreductase activity; IEA.
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                                                Similarity 6; Conserv
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                                                                                                                                                      proteome;
147 AA;
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                                                89.5%; ilarity 100.0%; Conservative
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100.0%; Pr/
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                                                     0
                                                                           Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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                                                                                                                                                      Oxidoreductase.
2745255C84FCCCCD CRC64;
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                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacillaceae; Bacillus;
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V; IEA.
V; IEA.
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                                                                           DB 2;
51;
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                                                                                                   Length 147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P., Han C., Hill I
Lucas S., Okinaka
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                                                     Indels
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RESULT 11
Q739P8 BACC1
ID Q739P8;
AC Q739P8;
DT 05-JUL-2004, integrated into UniProtKl
DT 05-JUL-2004, sequence version 1.
DT 07-MAR-2006, entry version 14.
DE General stress protein.
GN Name=dps; ORFNames=BCE 2002;
OS Bacillus cereus (strain ATCC 10987).
OC Bacteria; Firmicutes; Bacillales; Bac:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q813P2_BACCR
ID Q813P2_D
AC Q813P2_D
OT 01-JUN-2
DT 01
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Best Local S
Matches 6
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GO; GO:0003677; F:DNA binding; IEA.
GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0004322; F:ferroxidase activity; IEA.
GO; GO:0016491; F:oxidoreductase activity; IEA.
GO; GO:0016491; F:oxidoreductase activity; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006950; P:response to stress; IEA.
InterPro; IPR002177; DPS_DNA_bd.
InterPro; IPR002317; Ferritin_Dps.
InterPro; IPR002317; Ferritin_rel.
Pfam; PP00210; Ferritin_To.
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01-JUN-2003, integrated into UniProtKB/TrEMBL.
01-JUN-2003, sequence version 1.
07-FEB-2006, entry version 18.
Non-specific DNA-binding protein Dps / Iron-binding ferritin-like antioxidant protein / Ferroxidase (EC 1.16.3.1).
OrderedLocusNames=BC2011; ORFNames=BC 2011;
OrderedLocusNames=BC2011; ORFNames=BC 2011;
Bacillus cereus (strain ATCC 14579 / DSM 31).
Bacillus cereus group.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PR01346; HELNAPAPROT.
ProDom; PD149803; DPS; 1.
PROSITE; PS00818; DPS 1; 1.
Complete proteome; DNA-binding;
SEQUENCE 147 AA; 16680 MW; 3
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MEDLINE=22608415; PUMPdd=12721630; DOI=10.1038/nature01582;

Ivanova N., Sorokin A., Anderson I., Galleron N., Candelon B.,

Kapatral V., Bhattcacharyya A., Reznik G., Mikhailova N., Lapidus A.,

Chu L., Mazur M., Goltsman E., Larsen N., D'Souza M., Walunas T.,

Grechkin Y., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,

Overbeek R., Kyrpides N.C.;

"Genome sequence of Bacillus cereus and comparative analysis with

Bacillus anthracis.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HSSP; Q8RPQ1; 1JIG.
SMR; Q813P2; 2-147.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nature 423:87-91(2003).
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36A4D54D650CCCCD CRC64;
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            Bacillaceae;
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            Bacillus;
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Best Local
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GenomeReviews, AE017194 GR; dpS.
GO; GO:0008199; F:ferric iron binding; IE2
GO; GO:0006879; P:iron ion homeostasis; IIG
GO; GO:0006950; P:response to stress; IEA
InterPro; IPR002177; DPS DNA bd.
InterPro; IPR008331; Ferritin_Dps.
InterPro; IPR018347; Ferritin_rel.
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SMR; Q739P8; 2-147
MEDIINE=22608414; PubMed=12721629; DOI=10.1038/nature01586; Read T.D., Peterson S.N., Tourasse N.J., Baillie L.W., Paulsen I.V., Nelson K.E., Tettelin H., Fouts D.E., Eisen J.A., Gill S.R., Melson K.E., Tettelin H., Fouts D.E., Rilstone J., Wu M., Holtzapple E.K., Okstad O.A., Helgason E., Rilstone J., Wu M., Kolonay J.F., Beanan M.J., Dodson R.J., Brinkac L.M., Gwinn M.L., Beboy R.T., Madgu R., Daugherry S.C., Durkin A.S., Haft D.H., Nelson W.C., Peterson J.D., Pop M., Khouri H.M., Radune D., Benton J.L., Mahamoud Y., Jiang L., Hance I.R., Weldman J.F., Benton J.L., Mahamoud Y., Wolf A.M., Watkins K.L., Nierman W.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OBRPQ1 BACAN PRELIMINARY; PRT; 147 AA.
QBRPQ1; Q6HZV3; Q6KTT5; QB1RM9;
Q1-JUN-2002, integrated into UniProtKB/TrEMBL.
O1-JUN-2002, sequence version 1.
O7-MAR-2006, entry version 25.
Miniferritin Dlp2 (General stress protein).
Name=dlp2; Synonyms=dpS; OrderedLocusNames=BA2013,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PubMed=14960714; DOI=10.1093/nar/gkh258;
Rasko D.A., Ravel J., Oekstad O.A., Helgason E., Cer R.Z., Jia Shores K.A., Fouts D.E., Tourasse N.J., Angluoli S.V., Kolonay Nelson W.C., Kolstoe A.-B., Fraser C.M., Read T.D.;
"The genome sequence of Bacillus cercus ATCC 10987 reveals met adaptations and a large plasmid related to Bacillus anthracis Nucleic Acids Res. 32:977-988(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00210; Ferritin; 1.
PRINTS; PR01346; HELNAPAPROT.
ProDom; PD149803; DPS; 1.
PROSITE; PS00818; DPS_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.
MEDLINE=21964043; Pu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteria; Firmicutes;
Bacillus cereus group.
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NCBI_TaxID=222523;
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                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
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J. Biol. Chem. 277
                                                                                                                                                                                                                                                                                                                                                                                                          Papinutto
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                                                                                                                                                                                                                                                                                                                                                                                                          PubMed=11836250;
don W.G., Pitulis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89.5%; Score 34; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             iron binding; IEA. n homeostasis; IEA.
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N., Battistutta R., Montecu
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51;
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onay J.F.,
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RESULT 13
Q3EXQ4 BACTI
ID Q3EXQ4;
AC Q3EXQ4;
DT 08-NOV-2
DT 07-FEB--
DE NON-SPE
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GN ORFNAMES
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Matches 6
                                            QJEXQ4 BACTI PRELIMINARY; PRT; 154
QJEXQ4,
QJEXQ4,
QJEXQ4,
OB-NOV-2005, integrated into UniProtKB/T
OB-NOV-2005, sequence version 1.
O7-FEB-2006, entry version 3.
Non-specific DNA-binding protein Dps / I
antioxidant protein / Ferroxidase CEC 1.
ORFNames=RBTH 06140;
Bacillus thuringiensis serovar israelens
    Bacteria, Firmicutes, Bacillales, Bacillaceae, Bacillus cereus group.
NCBI TaxID=339854;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PDB; 1JIG; X-ray; A/B/C/D2-147.
GenomeReviews; AE016879_GR; BA2013.
GenomeReviews; AE017225_GR; BAS1871.
GenomeReviews; AE017334_GR; GBAA2013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Complete genome sequence of Bacillus anthracis Submitted (JAN-2004) to the EMBL/GenBank/DDBJ da
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brettin T.S., Bruce D., Challacom
Hitchcock P., Jackson P., Keim P.
Richardson P., Rubin E., Tice H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (MAY-2004) to
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Fraser C.M.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "The genome sequence of Bacillus anthracis closely related bacteria.";
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                                                                                                                                                                                                                                                                                                 SEQUENCE 147 AA
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ProDom; PD149803; DPS; 1.
PROSITE; PS00818; DPS_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Copyrighted by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=Sterne;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Bacillus anthracis comparative genomics."; Submitted (MAY-2004) to the EMBL/GenBank/DE
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                                                                                                                                                                                                                                                                                                                                                            Pfam; PF00210; Ferritin;
                                                                                                                                                                                                                                                                                                                                                                   GO; GO:0006879; P:iron ion homeosta
GO; GO:0006950; P:response to streat
InterPro; IPR002177; DPS DNA bd.
InterPro; IPR008331; Ferritin Dps.
InterPro; IPR012347; Ferritin_rel.
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GO:0006879;
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AE016879; AAP25903.1; -;
AE017225; AAT54186.1; -;
AE017334; AAT31135.1; -;
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                                                                                                                                                                                                                                                 Similarity 100.0%; 6; Conservative
                                                                                                                                                                                                    WMLSAF 145
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                                                                                                                                                                                                                                                                          89.5%;
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                                                                                                                                                                                                                                                                                                 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                      homeostasis;
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Genomic_DNA.
Genomic_DNA.
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                                                                      / Iron-binding ferritin-like 1.16.3.1).
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51;
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re J.,
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databases.
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Koehler T.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              databases
                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                          Length 147;
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                                      Bacillus
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s., Okinaka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Federova
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Best Local
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GG; GC:0003677; F:DAA binding; IEA.
GG; GC:00008199; F:ferric iron binding
GG; GC:00008192; F:ferroxidase activi
GG; GC:0004322; F:ferroxidoreductase activi
GG; GC:0006491; F:oxidoreductase activi
GG; GC:0006879; P:iron ion homeosetas;
GG; GC:0006879; P:response to stress
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                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
PubMed=15703394; DOI=10.1073/pnas.0409900102;
Ruby E.G., Urbanowski M., Campbell J., Dunn A., Faini M.,
Lostroh P., Lupp C., McCann J., Millikan D., Schaefer A.,
Stevens A., Visick K., Whistler C., Greenberg E.P.;
"Complete genome sequence of Vibrio fischeri: a symbiotic
with pathogenic congeners.";
Proc. Natl. Acad. Sci. U.S.A. 102:3004-3009(2005).
EMBL; CP000020; AAW85552.1; -; Genomic_DNA. GO; GO:0016020; C:membrane; IBA. GO; GO:0005293; F:lysine permease activity; GO; GO:0006865; P:amino acid transport; IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OSES24,

IS-MAR-2005, integrated into I

15-MAR-2005, sequence version

07-FEB-2006, entry version 5.
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InterPro; IPR012347; Ferritin_rel.
Pfam; PF00210; Ferritin; 1.
                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprodistributed under the Creative Commons Attribution-NoDerivs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria;
Vibrionaceae; Vibrio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vibrio fischeri (strain ATCC 700601 / ES114).
Bacteria; Proteobacteria; Gammaproteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lysine exporter protein. OrderedLocusNames=VF1057;
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Probom; PD149803; DPS; 1.
PROSITE; PS00818; DPS 1; 1.
DNA-binding; Oxidoreductase.
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GO:0008199; F:ferric iron binding; IEA
GO:0004322; F:ferroxidase activity; IE
GO:0016491; F:oxidoreductase activity;
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53;
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Stabb E.,
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Matches 6
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Best Local
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Pfam; PF01810; LysE; 1
Complete proteome.
SEQUENCE 211 AA; 22
                                                                                                                                                                                                                                                                       10-JAN-2006, integrated
10-JAN-2006, sequence v
07-MAR-2006, entry vers
Permease of the major f
    "Complete
Bacterium
                                                                                                                                                                                        Magnetospirillum magneticum AMB-1.
Bacteria; Proteobacteria; Alphaproteobacteria;
Rhodospirillaceae; Magnetospirillum.
                                                                                                                                                                                                                                                                                                                                                            QZW9C2_MAGSA
Q2W9C2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Copyrighted by the UniProt Consortium, see http://www.uniprotorty.to
                                                                                     PubMed=16303747;
                                                                                                         STRAIN=AMB-1
                                                                                                                                                                  NCBI_TaxID=342108;
                                                                                                                                                                                                                                                          ORFNames=amb0749;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
                                           Takeyama H.;
                                                             Matsunaga T., Okamura
                                                                                                                             NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            138
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Genome Sequence of Magnetospirillum sp
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                                                               Fukuda Y.,
  sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 34;
Pred. No.
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  the Facultative A
p. strain AMB-1.";
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Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                     PRT;
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                                                                                                                                                                                                                                                                             superfamily
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                                                               Wahyudi A.T.,
                                                                                                                                                                                                                                                                                                                                                                                     458
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73;
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                                                                                                                                                                                                                Rhodospirillales;
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                                                               Murase Y.,
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RESULT 18
p88944 HHV8
ID p88944;
DT 01-MAY-1997,
DT 01-MAY-1997,
DT 07-FEB-2006,
DE ORF 56 (DNA 1
OS Human herpes
OC Viruses; dsbb
OC Gammaherpesvicox
OX NCBI_TaxID=37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q44233
ID 44233
ID 7010
DT 011
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Best Local
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Best Local
P88944;
01-MAY-1997, integrated into UniProtKB/T
01-MAY-1997, sequence version 1.
07-FEB-2006, entry version 23.
ORF 56 (DNA replication protein).
Human herpesvirus 8 (HHV-8) (Kaposi's saviruses; dsDNA viruses, no RNA stage; He Gammaherpesvirinae; Rhadinovirus.
NCBI_TaxID=37296;
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Q44233
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GO; GO:0009341; C:beta-galactosidase complex; IEA.
GO; GO:0004565; F:beta-galactosidase activity; IEA.
GO; GO:0005975; P:carbohydrate metabolism; IEA.
InterPro; IPR003476; Glyco-hydro_42.
Pfam; PF02449; Glyco-hydro_42; 1.
PIRSF; PIRSF001084; B-galactosidase; 1.
SEQUENCE 637 AA; 70895 MW; 98160B4FCC3A728C CRC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
MEDLINE=95238267; PubMed=7721689;
MEDLINE=95238267; PubMed=7721689;
Gutshall K.R., Trimbur D.E., Kasmir J.J., Brenchley J.E.;
"Analysis of a novel gene and beta-galactosidase isozyme i
psychrotrophic Arthrobacter isolate.";
J. Bacteriol. 177:1981-1988(1995).
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07-FEB-2006, entry version 20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Copyrighted
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SUBCELLULAR LOCATION:
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AA; 48181
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 34; DB 2;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                     PRT;
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                                                  Herpesviridae;
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Attribution-NoDerivs License
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1.6e+02;
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RESULT 19
Q3WYR2 9ACTN
ID Q3WYR2;
AC Q3WYR2;
DT 11-OCT-2005, 8
DT 07-MAR 2006, 9
DT 07-MAR 2006, 9
DT VA1yl-tRNA 8yr
GN ORFNAMESERY!I
OS Rubrobacter x
OC Rubrobacter x
OC Rubrobacterin Active
CR NUCBI_TAXID=266
RN [1]
RP NUCLEOTIDE SER
RC STRAIN=DSM 99
RG US DOB Joint G
RA Copeland A. 1
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Best Local
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Parry J.P., l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U75698; AAC57138.1; -; Genomic_DNA.
EMBL; U93872; AAB82614.1; -; Genomic_DNA.
GO; GO:0003896; F:DNA primase activity; IEA.
GO; GO:0006260; P:DNA replication; IEA.
InterPro; IPR004340; UL52 UL70.
Pfam; PF03121; Herpes_UL52; 1.
SHOULENCE 843 AA; 95554 MW; 89248BFCB7DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neipel F., Albrecht J.-C., Ensser A., Huang Y. Friedman-Kien A.E., Fleckenstein B.; "The genome of human herpervirus 8 cloned from Submitted (JUL-2001) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WEDLING=7/2/2... Fleckenstein Weipel F. Albrecht J.-C., F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Russo J.J., Bohenzky R.F. Parry J.P., Peruzzi D., Submitted (MAY-1997) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genes by KSHV."
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     US DOE Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
                                                                                                                                                                                                                 ORFNames=RxylDRAFT_0191;
Rubrobacter xylanophilus DSM 9941.
                                                                                                                                                                                                                                                                                     11-OCT-2005, integrated into t
11-OCT-2005, sequence version
07-MAR-2006, entry version 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Science 274:1739-1744(1996)
                                                                                                                                          Rubrobacterineae; |
NCBI_TaxID=266117;
                                                                                                                                                                           Bacteria; Actinobacteria; Rubrobacteridae; Rubrobacterales;
Rubrobacterineae; Rubrobacteraceae; Rubrobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=97296220; PubMed=9151804;
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                                                                           STRAIN=DSM 9941;
                                                                                                  NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                 Valyl-tRNA synthetase,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Distributed
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rol. 71:4187-4192(1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bohenzky R.A., Chien M.-C., Chen J., Yan M., Maddalen Peruzzi D., Edelman I.S., Chang Y., Moore P.S.; sequence of the Kaposi sarcoma-associated herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Acad. Sci.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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Pred. No.
                                                                                                                                                                                                                                                                                                                                          UniProtKB/TrEMBL
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3e+02;
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       Detter P.;
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RESULT 20
Q73RB9_TREDE
ID Q73RB9_TREDE
AC Q73RB9;
DT 05-UUL-2004, i
DT 05-UUL-2004, i
DT 07-MAR-2006, e
DT 07-MAR-2006, e
DT Nuclease SbcCl
GN OrderedLocusNa
OS Treponema dent
OC Bacteria, Spin
OX NCBI_TaxID=158
RN [1]
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                                                                                                                                                                                          05-JUL-2004, sequence version 07-MAR-2006, entry version 20. Nuclease SbCCD, C subunit, put OrderedLocusNames=TDE0171; ORF
NUCLEOTIDE SEQUENCE (LARGE SCALE GENOMIC DNA)
STRAIN-ATCC 35405 / DSM 14222;
PubMeda-15064399; DOI=10.1073/ppnas.0307639101;
Seshadri R., Myers G.S.A., Tettelin H., Eisen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00133; tRNA-synt 1; 1.

PRINTS; PR00986; TRNASYNTHAL.

TIGR0422; vals; 1.

PROSITE; PS00178; AA_TRNA_LIGASE_I; 1.

PROSITE; PS00178; AA_TRNA_LIGASE_I; 1.

ATP-binding; Aminoacy1-tRNA_synthetase; Ligase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copeland A., Lucas S., Lapidus A., Barry K., Detter C., Hammon N., Israni S., Pitluck S., Richardson P.; Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Sequencing of the draft genome and xylanophilus DSM 9941.";
Submitted (JUL-2005) to the EMBL/Ger
                                                                                                                                                  Bacteria; Spirochaetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR002300; tRNA-synt_1a.
InterPro; IPR001412; tRNA-synt_I.
InterPro; IPR002303; tRNA-synt_val.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/teDistributed under the Creative Commons Attribution-NoDerivs License
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                                                                                                                 NCBI_TaxID=158;
                                                                                                                                                                         Treponema denticola.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO:0016874; F:ligase activity; IEA.
GO:0000166; F:nucleotide binding; IEA.
GO:0004832; F:valine-tRNA ligase activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preliminary data.
CATALYTIC ACTIVITY: ATP + L-valine + tRNA(Val)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9941."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        607
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UBUNIT: Monomer (By similarity).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          biosynthesis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WILSAFS 613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Lucas S., Lapidus A., Barry K.,
srani S., Pitluck S., Richardson
                                                                                                                                                                                                                                                                                               integrated into UniProtKB/TrEMBL
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                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97415 MW;
                                                                         [LARGE SCALE GENOMIC DNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89.5%;
85.7%;
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                                                                                                                                               Spirochaetales; Spirochaetaceae;
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                                                                                                                                                                                             putative.
ORFNames=TDE_0171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 34; DB Pred. No. 3.1e.
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                               1030 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Db ...
3.1e+02;
0;
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     J.A.,
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RESULT 21
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ID Q4B83
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Q4B833;
13-SEP-2005,
13-SEP-2005,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dodson R.J., Davidsen T.M., DeBoy R.T., Fouts D.B., Haft D.H., Selengut J., Ren Q., Brinkac L.M., Madupu R., Kolonay J.F., Durkin S.A., Daugherty S.C., Shetty J., Shvartsbeyn A., Gebregeorgis E., Geer K., Tsegaye G., Malek J.A., Ayodeji B., Shatsman S., McLeod M.P., Smajs D., Howell J.K., Pal S., Amin A., Vashisth P., McNeill T.Z., Xiang Q., Sodergren E., Baca E., Weinstock G.M., Norris S.J., Fraser C.M., Paulsen I.T.; "Comparison of the genome of the oral pathogen Treponema denticola
                                                                Copeland A., Lucas S., Lapidus A., Barry K., Detter C., Hammon N., Israni S., Pitluck S., Richardson P.; Submitted (JUN-2005) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Copeland A., Lucas S., Lapidus A., Barry K., Hammon N., Israni S., Pitluck S., Richardson "Sequencing of the draft genome and assembly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                             US DOE Joint Copeland A.,
                                                                                                                                                                    STRAIN=G4;
                                                                                                                                                                                                                                       Submitted
                                                                                                                                                                                                                                                                                                                                    STRAIN=G4;
US DOE Joint Genome Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Sequencing of the draft vietnamiensis G4.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US DOE Joint
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacteria; Proteobacteria; Betaproteobacteria;
Burkholderiaceae; Burkholderia; Burkholderia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Burkholderia vietnamiensis G4.
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GO; GO:0016887; F:ATPase activity;
InterPro; IPR003439; ABC_transp_lik
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GenomeReviews; AE017226_GR; TDE0171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       with other spirochete genomes.";
Proc. Natl. Acad. Sci. U.S.A. 101:5646-5651(2004).
                                                                                                                                                                                                                                                                                       "Annotation
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                 CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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                                                                                                                Genome Institute (JGI-PGF);
Lucas S., Lapidus A., Barry K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry version
                                                                                                           Lucas S.,
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                                                                                                                                                                                                                                                                                                                                      (JGI-ORNL)
                                                                                                                                                                                                                                                                                    assembly of Burkholderia vietnamiensis
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cepacia complex.
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RESULT 23
Q2UDO8 ASPOR
ID QZUDO8;
ID QZUDO8;
DT 24-JAN-2006, s
DT 07-MAR-2006, s
DT 07-MAR-2006, s
DT 07-MAR-2005, s
DT 07-MAR-2007
GN ORFNames=A0090
OS Aspergillus or
OC Eukaryota; Fun
OC Eukaryota; Fun
OC Eurotiales; Tx
OX NCBI TaxID=506
RN [1]
RP NUCLEOTIDE SEQ
RC STRAIN=RIB 40;
RX PubMed=1637201
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GO; GO:00
Pyruvate.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Liu B., Zhao B., Wang X.Y., Xu Y.Y., Liu Y.Q., Song L., Ye Sheng H., Gao Y., Zhang C.L., Wei Y.J., Zhang J., Song L., Zhao Z.W., Ding J.F., Liu L.S., Gao R.L., Wu Q.Y., Qiang B. Yuan J.G., Liew C.C., Zhao M.S., Hui R.T.; Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q5QTR8;
04-JAN-2005, integrated into U
04-JAN-2005, sequence version
07-FEB-2006, entry version 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                             24-JAN-2006, integrated into UniProtKB/TrEMBL. 24-JAN-2006, sequence version 1. 07-MAR-2006, entry version 3. Hydrolases of the alpha/beta superfamily.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
NUCLEOTIDE SEQUENCE.
STRAIN=RIB 40;
PubMed=16372010; DOI=10.1038/nature04300;
                                                                     Aspergillus oryzae.
Eukaryota; Fungi; Ascomycot:
Eurotiales; Trichocomaceae;
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                                 ORFNames=A0090012000367;
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HUMAN
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les 6; Conserv
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GO:0016491; F:oxidoreductase activity; IEA.
GO:0006118; P:electron transport; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                 AF109190; AAQ13500.1; -; mRNA.
NCE 44 AA; 5210 MW; 90754272D3B001A4 CRC64;
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                                                                                        Ascomycota;
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100.0%; Prr
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                                                                      a; Pezizomycotina; Eurotiomycetes; mitosporic Trichocomaceae; Aspergillus.
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Pred. No.
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Pred. No. 24;
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hes 0;
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X Kusumoto K., Arima T., Akita O., Kashiwagi Y., Abe K., Gomi K.,

XA Horiuchi H., Kitamoto K., Kobayashi T., Takeuchi M., Denning D.W.,

XA Galagan J.E., Nierman W.C., Yu J., Archer D.B., Bennett J.W.,

Bhatnagar D., Cleveland T.E., Fedorova N.D., Gotoh O., Horikawa H.,

Ab Hosoyama A., Ichinomiya M., Igarashi R., Iwashita K., Juvvadi P.R.,

XA Kato M., Kato Y., Kin T., Kokubun A., Maeda H., Maeyama N.,

XA Kato M., Kato Y., Kin T., Kokubun A., Maeda H., Maeyama N.,

XA Maruyama J., Nagasaki H., Nakajima T., Oda K., Okada K., Paulsen I.,

XA Maruyama J., Sawano T., Takahashi M., Takase K., Terabayashi Y.,

XA Wortman J.R., Yamada O., Yamagata Y., Anazawa H., Hata Y., Koide Y.,

XA Komori T., Koyama Y., Minetoki T., Suharnan S., Tanaka A., Isono K.,

XI "Genome sequencing and analysis of Aspergillus oryzae.";

Nature 438:1157-1161(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                               PubMed=14702039, DOI=10.1038/ng1285;
PubMed=14702039, DOI=10.1038/ng1285;
Ota T., Suzuki Y., Nishikawa T., Otsuki T., Sugiyama T., Irie R.,
Wakamatsu A., Hayashi K., Sato H., Nagai K., Kimura K., Makita H.,
Sekine M., Obayashi M., Nishi T., Shibahara T., Tanaka T., Ishii S.,
Yamamoto J., Saito K., Kawai Y., Isono Y., Nakamura Y., Nagahari K.,
Murakami K., Yasuda T., Iwayanagi T., Wagatsuma M., Shiratori A.,
Sudo H., Hosoiri T., Kaku Y., Kodaira H., Kondo H., Sugawara M.,
Takahashi M., Kanda K., Yokoi T., Furuya T., Kikkawa E., Omura Y.,
Abe K., Kamihara K., Katsuta N., Sato K., Tanikawa M., Yamasaki M.,
Ninomiya K., Ishibashi T., Yamashita H., Murakawa K., Fujimori K.,
Ninomiya K., Ishibashi T., Yamashita H., Murakawa K., Fujimori K.,
Tanai H., Kimata M., Watanabe M., Hiraoka S., Chiba Y., Ishida S.,
Ono Y., Takiguchi S., Watanabe S., Yosida M., Hotuta T., Kusano J.,
Ono Y., Takiguchi S., Watanabe S., Yosida M., Hotuta T., Kusano J.,
Ono Y., Takiguchi S., Watanabe S., Yosida M., Hotuta T., Kusano J.,
Ono Y., Takiguchi S., Watanabe S., Yosida M., Hotuta T., Nomura Y.,
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LHUMAN
PRG4 HUMAN STANDARD; PRT; 1404 AA.

[92954; Q6DNC4; Q6DNC5; Q6ZMZ5; Q9BX49;
06-DEC-2005, integrated into UniProtKB/Swiss-Prot.
06-DEC-2005, sequence version 2.
07-MAR-2006, entry version 38.

Proteoglycan-4 precursor (Lubricin) (Megakaryocyte stimulating factor)
(Superficial zone proteoglycan) [Contains: Proteoglycan-4 C-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                         NÜCLEOTIDE SEQUENCE [MRNA] (ISOFORM A), AND VARIANT TRP-180. Turner K.J., Fitz L.J., Temple P., Jacobs K., Larson D., Leary Kelleher K., Giannotti J., Calvetti J., Fitzgerald M., Kriz M. Ferenz C., Grobholz J., Fraser H., Bean K., Norton C.R., Gesne Bhatla S., Kriz R., Hewick R., Clark S.C.; Submitted (SEP-1996) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=PRG4; Synonyms=MSF, SZP;
Homo sapiens (Human)
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Euarchont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                         TISSUE=Synovial cell
                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA] (ISOFORM E).
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GO:0016787;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BAE60557.1; -; Genomic DNA. F:hydrolase activity; TEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         33626 MW;
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85.7%;
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Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 304;
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VA Yoshikawa Y., Matsunawa H., Ichihara T., Shiohata N., Sano S., Momiyama H., Satoh N., Takamai S., Terashima Y., Suzuki O., A Makagawa S., Senoh A., Mizoguchi H., Goto Y., Shimizu F., Wakebe H., A Hishigaki H., Watanabe T., Sugiyama A., Takemoto M., Kawakami B., Yamazaki M., Watanabe T., Sugiyama A., Takemoto M., Kawakami B., Yamazaki M., Watanabe K., Kumagai A., Itakura S., Fukuzumi Y., A Yamazaki M., Watanabe K., Kumagai A., Tanigami A., Fujiwara T., Ono T., Yamada K., Fujii Y., Ozaki K., Hirao M., Ohmori Y., Okamoto S., Ohitani R., Kawakami T., Kobatake N., Inagaki H., Ikema Y., Okamoto S., Okitani R., Kawakami T., Noguchi S., Itoh T., Shigeta K., Senba T., A Matsumura K., Nakajima Y., Mizuno T., Morinaga M., Sagaki M., A Togashi T., Oyama M., Hata H., Watanabe M., Komatsu T., Nakagawa K., A Mizushima-Sugano J., Satoh T., Shirai Y., Takahashi Y., Nakagawa K., A Makai K., Yada T., Namura N., Kikuchi H., Masuho Y., Yamashita R., Nakai K., Yada T., Nakamura Y., Ohara O., Isogai T., Sugano S.; "Complete sequencing and characterization of 21,243 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABDLINE=99120896; running factor precursor and is a multifunctional proteoglycan with potential growth-promoting, cytoprotective, and proteoglycan with potential growth-promoting, cytoprotective, and proteoglycan with potential growth-promoting.

"Articular cartilage superficial zone protein (SZP) is homologous commegakaryocyte stimulating factor precursor and is a multifunctional megakaryocyte stimulating factor precursor and cytoprotective, and proteoglycan with potential growth-promoting, cytoprotective, and proteoglycan with potential growth-promoting."

"Traction, AND "Traction, 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PubMed=14976050; DOI=10.1182/blood-2003-06-1825; Liu Y.J., Lu S.H., Xu B., Yang R.C., Ren Q., Liu Yan F.Y., Han Z.B., Han Z.C.;
GLYCOSYLATION.
MEDLINE=94241694;
Schumacher B.L.,
                                                                                                                                                         "A comparison of vitronectin and (In) Preissner K.T., Rosenblatt Mosher D.F. (eds.);
                                                                                                                                                                                                                              Merberg D.M., Fitz L.J., Temple P., Giannotti J., Mu
Fitzgerald M., Scaltreto H., Kelleher K., Preissner
Jacobs K., Turner K.;
                                                                                                                                                                                                                                                                                                                                                                        stimulating activity.";
Blood 78:279A-279A(1991).
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SUBCELLULAR LOCATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jones A.R., Hughes C.E., Flannery C.R., Caterson B.; "Cloning and production of recombinant PRG4/cartilage superficial zone proteoglycan (SZP) N- and C-terminal domains."; Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Urine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Hemangiopoietin,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ngiopoietin, a novel human growth factor for the th hematopoietic and endothelial cell lineages."; 103:4449-4456(2004).
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                                                                                                  of vitronectins
Publishers B.V.,
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PubMed=8185311;
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                                                                                                                                                                                  megakaryocyte stimulating S., Kost C., Wegerhoff J.,
DOI=10.1006/abbi.1994.1219; id T.M., Aydelotte M.B.,
                                                                                                        receptors,
n (1993).
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Marcelino U.,
Thing C., 5
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Robbins C., Sood R., Makalowska I., Baxevanis A., Johnstone
Laxer R.M., Zemel L., Kim C.A., Herd J.K., Ihle J., Williams
Johnson M., Raman V., Alonso L.G., Brunoni D., Gerstein A.,
Papadopoulos N., Bahabri S.A., Trent J.M., Warman M.L.;
"CACP, encoding a secreted proteoglycan, is mutated in campt
arthropathy-coxa vara-pericarditis syndrome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jay G.D., Britt D.E., Cha C.-J.;
"Lubricin is a product of megakaryocyte stimulating expression by human synovial fibroblasts.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "A novel proteoglycan superficial zone of ar
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J. Proteome Res. 4:2070-2080(2005)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu T., Qian W.-J., Gritsenko Moore R.J., Smith R.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=10743795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (proteoglycan 4) genes.
Cytogenet. Cell Genet.
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"Isolation, characterization and mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=10545950; DOI=10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Arch. Biochem. Biophys.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Human plasma N-glycoproteome analysis by immunoaffinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=20573856;
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                                                                                                                                               Note=No experiments.

Name=F; Synonyms=Hemangiopoietin, HAPO;
IsoId=092954-6; Sequence=VSP 016469;
ISOId=092954-6; Sequence=VSP 016469;
TISSUE SPECIFICITY: Highly expressed in synovial tissue, cartilage and liver, and weakly in heart and lung. Isoform B is expressed in liver, and weakly in heart and brain. Isoform C and isoform D are
                                                          kidney, lung, liver, heart and brain. Isoform widely expressed.
PTM: N-glycosylated.
PTM: O-glycosylated; contains glycosaminoglyc sulfate and keratan sulfate.
PTM: The disulfide bond between Cys-1146 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBUNIT: Homodimer; disulfide-
SUBCELLULAR LOCATION: Secreted
ALTERNATIVE PRODUCTS:
for protein cleavage (By similarity). DISEASE: Defects in PRG4 are the cause of camptodactyly-arthropathy-coxa vara-pericarditis syndrome (CACP) [MIM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 surface. FUNCTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      articulating joints. Prevents protein deposition onto cartilage from synovial fluid, control adhesion-dependent synovial growth and inhibit the adhesion of synovial cells to the cartilage
                                                                                                                                                                                                                                                                                                                                                                                                      Name=C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Event=Alternative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primitive cells
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                                                                                                                                                                                                                                                                        experimental confirmation available;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Plays a role for boundary lubrication within
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AND INVOLVEMENT
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of both h
                                                                                                                                                                                                                                                                                            Sequence=VSP_016467, VSP_016470,
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                                                                                                                                                                                                                                                                                                                                   Sequence=VSP_016467, VSP_016468;
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[MIM:208250];

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Best Local
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13-SEP-2005,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                 Complete
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ralstonia eutropha (strain JMP134) (Alcaligenes eutrophus).
Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
Burkholderiaceae; Cupriavidus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL;
19-JUL-2005, integrated into UniProtKB/TrEMBL.
19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 3.
Chromosome 7 SCAF14703, whole genome shotgun in Grenames=GSTENG00022129001;
Tetraodon nigroviridis (Green puffer).
Eukaryota; Metazoa; Chordata; Craniata; Vertel
                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLECTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C., Glavina Hammon N., Israni S., Pitluck S., Goltsman E., Martinez M., Schmutz J., Larimer F., Land M., Lykidis A., Richardson P.;
"Complete sequence of chromosome 2 of Ralstonia eutropha JMP134.";
Submitted (AUG-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q46TT1_RALEJ
Q46TT1;
                                                                                                      Q4S8Z0_TETNG
Q4S8Z0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OrderedLocusNames=Reut_B4097;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    + +
                                                                                                                                                                                                                                                                                                                                             EMBL; CP000091; AAZ63453.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                         Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                         Copyrighted
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RALEJ
                                                                                                                                     TETNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       also called Jacobs syndrome. CACP is an autosomal recession disorder. Individuals with CACP have normal appearing joints at birth but with advancing age develop joint failure associated with noninflammatory synoviocyte hyperplasia and subintimal fibrosis of the synovial capsule.

MISCELLANEOUS: Different forms of various molecular weight have miscellaneous are possibly due to different levels of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   been observed. Such forms are possibly due to diffe glycosylation and protein cleavage (By similarity). SIMILARITY: Contains 2 hemopexin-like domains. SIMILARITY: Contains 2 SMB. (somatomedin B) domains.
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ilarity 71.4%;
Conservative
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8424 MW; 39
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                                                                                                                                                                                                                                                     ; Score 32; DB;
; Pred. No. 67;
2; Mismatches
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Pred. No. 8.4e
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   Vertebrata; Euteleostomi;
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Q43Q96_SOLUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                      13-SEP-2005, 13-SEP-2005, 1
                                                                                                                          US DOB Joint Genome Institute (JGI-PGF);
Copeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
"Sequencing of the draft genome and assembly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
Genoscope; Whitchead Institute Centre for Genome Researc Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                                                                                                                                                                                                                                                                                                                                                                                     Q43Q96_SOLUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL;
US DOE Joint Genome Institute (JGI-ORNL);
Larimer F., Land M.;
"Annotation of the draft genome assembly
                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
STRAIN=Ellin6076;
                                                                                                                                                                                                                                                                                                         ORFNames=AcidDRAFT_11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Copyrighted by the UniProt Consortium, Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Acanthomorpha; Acanthopterygii; Tetradontoidea; Tetradontidae;
                                                STRAIN=Ellin6076;
                                                                                              Submitted (JUN-2005)
                                                                                                             "Sequencing of the Ellin6076.";
                                                                                                                                                                                                                                                                       Bacteria; Acidobacteria;
                                                                                                                                                                                                                                                                                     Solibacter usitatus Ellin6076
                                                                                                                                                                                                                                                                                                                     Transport-associated
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                                                              NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                      NCBI_TaxID=234267;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  preliminary data.
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NCE 117 AA; 12900 MW; F
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                                                                                                                                                                                                                                                          Solibacter.
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71.4%;
                                                                                                                                                                                                                                                                                                                      precursor.
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Pred. No. 1.1e+02;
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Tetraodon.
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Attribution-NoDerivs License
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ID Q9WXJ3 PORGI
ID Q9WXJ3 PAR
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Q9WXJ3;
01-NOV-1999,
01-NOV-1999,
07-FEB-2006,
                                                                                                                                                                     Prodom; P
PROSITE;
PROSITE;
SEQUENCE
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SIGNAL
                                                                                                                                                                                                                                                                                   GO; GO:0008199; F:ferric iron binding; IEA.
GO; GO:0006879; P:iron ion homeostasis; IEA.
GO; GO:0006870; P:response to stress; IEA.
InterPro; IPR002177; DPS DNA bd.
InterPro; IPR008331; Ferritin Dps.
InterPro; IPR012347; Ferritin Tel.
Pfam; PF00210; Ferritin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=ATCC 33277;
MEDLINE=22483669; PubMed=12595429;
DOI=10.1128/IA1.71.3.1170-1178.2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AB025779; BAA76886.1; -; Genomic_DNA. HSSP; Q8RPQ1; 1JIG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Purification, gene cloning, gene expression, and the obligate anaerobe Porphyromonas gingivalis."; Infect. Immun. 71:1170-1178(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nakayama K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ueshima J., Shoji M., Ratnayake D.B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Porphyromonadaceae; Porphyromonas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Porphyromonas gingivalis (Bacteroides gingivalis).
Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteroidales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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PROSITE; PS50914; BON; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro;
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                                                                                            Local
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                                                                                                                                                                  3; PR01346; HELNAPAPROT.
1; PD149803; DPS; 1.
1%; PS00818; DPS 1; 1.
1%; PS00819; DPS 2; 1.
1CE 159 AA; 17793 MW;
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                                                                           Conservative
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Pred. No.
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                                                                                                 1.5e+02;
                                                                                                                        DB 2;
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     OCCUPED DD AND
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ID Q84S36_ORYSA
AC Q84S36;
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Best Local S
Matches 5
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Q7MXS1;
15-DEC-2003,
15-DEC-2003,
07-FEB-2006,
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01-JUN-2003,
07-FEB-2006,
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TIGR; PG0090; -
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SEQUENCE 159 AA
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5; Conser
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Hypothetical protein 0J9990_A01.131 (Hypothet 0J1449 C01.14).

Name=0J9990_A01.131; Synonyms=0J1449 C01.14; Oryza sativa (japonica cultivar-group). Embr. Eukaryota; Viridiplantae; Streptophyta; Embr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ProDom; PD149803; DPS; 1.
PROSITE; PS00818; DPS_1; 1.
PROSITE; PS00819; DPS_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; PF00210; Ferritin;
PRINTS; PR01346; HELNAPAI
ProDom; PD149803; DPS; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Complete genome sequence of the oral pathogenic bacterium Porphyromonas gingivalis strain W83.";
J. Bacteriol. 185:5591-5601(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOI=10.1128/JB.185.18.5591-5601.2003;
Nelson K.E., Fleischmann R.D., DeBoy R.T., Paulsen I.T., Fouts Eisen J.A., Daugherty S.C., Dodson R.J., Durkin A.S., Gwinn M. Eisen J.A., Kolonay J.F., Nelson W.C., Mason T.M., Tallon L., G. Granger D., Tettelin H., Dong H., Galvin J.L., Duncan M.J., Dewhirst F.E., Fraser C.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OrderedLocusNames=PG0090; ORFNames=PG_0090;
Porphyromonas gingivalis (Bacteroides gingivalis).
Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteria; Porphyromonadaceae; Porphyromonas.
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OrderedLocusNames=PG0090;
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GO; GO:0008199; F:ferric iron binding; IEA
GO; GO:0006879; P:iron ion homeostasis; IEA
GO; GO:0006950; P:response to stress; IEA.
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InterPro; IPR008331; Ferr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA]
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                                                                                                                                    entry version 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17865 MW;
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Ferritin_Dps.
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                                                                                                                                                                            into UniProtKB/TrEMBL
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ITIVAT-group).
Streptophyta; Embryophyta; Tracheophyta;
yta; Liliopsida; Poales; Poaceae; BEP clade;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UniProtKB/TrEMBL
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60889B904D331AC6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                           ed. No. 1.5e+02;
Mismatches 0
                                                                                                          (Hypothetical
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RASULT 31
Q4SFB2 TETING
ID Q4SFB2 T
D7 19-JUL-2
D8 Chromoso
ON Tetradon
ON OCT Tetradon
ON NCBI-Tax
RN [1]
RP NUCLEOTI
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RA Kellis M
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                                                                                                                                                                                                                                                                  RX PubMed=15496914; DOI-10.1038/nature03025;
RA Jaillon O., Aury J.-M., Brunet F., Petit J.-L., Stange-Thomann N.,
RA Jaillon O., Aury J.-M., Brunet F., Petit J.-L., Stange-Thomann N.,
RA Mauceli E., Bouneau L., Fischer C., Oouf-Costaz C., Bernot A.,
RA Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
RA Anthouard V., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
RA Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,
RA Cruaud C., Duprat S., Brottier P., Coutanceau J.-P., Gouzy J.,
RA Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
RA Farra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
RA Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
RA Laudet V., Schachter V., Quetier F., Saurin W., Scarpelli C.,
RA Wincker P., Lander E.S., Weissenbach J., Roest Crollius H.;
"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
RT The early vertebrate proto-karyotype.";
RN Nature 431:946-957(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                         Tetraodon nigroviridis (Green puffer).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q4SFB2;
19-JUL-2005, integrated into UniProtKB/TrEMBL.
19-JUL-2005, sequence version 1.
07-FEB-2006, entry version 3.
Chromosome 6 SCAF14605, whole genome shotgun sequence.
ORFNames=GSTENG00019173001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical SEQUENCE 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sasaki T., Matsumoto T., Katayose Y.; "Oryza sativa nipponbare(GA3) genomic DNA, clone:OJ9990_A01.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q4SFB2_TETNG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (A\overline{U}G-2002) to the EMBL/GenBank/DDBJ databases.
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NCBI_TaxID=39947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gramene; Q84S36;
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; AP005605; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WMLSAFS 7
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167 AA; 1
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        data
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Pred. No. 1.6e+02;
2; Mismatches 0
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genomic
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RESULT 32
PUT2_FUGRU
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01-AUG-1998,
07-FEB-2006,
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SEQUENCE
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RIBOLIDI Tunnicliffe G.R., Platzer M., Nyakatura G., Elgi
Brenner S., Rosenthal A.;
"Analysis of the genomic loci of Fugu rubripes homologs
"Analysis of the genomic loci of Fugu rubripes homologs
disease genes LiCAM, G6PD and P55.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases
-!- SUBCELLULAR LOCATION: Membrane; multi-pass membrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-2006, entry version 28.
Putative protein 2 (PUT2) (Fragment).
Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Neopterygii; Teleosteei; Euteleostei; Neoteleostei;
Acatihopterygii, Reanthopterygii; Percomorpha; Tetradontiformes;
Tetradontoidea; Tetraodontidae; Takifugu.
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                                                                                                                                                                                                                                                                                                                                               InterPro; IPR004153; CXCXC_repeat.
InterPro; IPR008853; TMEM9;
PANTHER; PTHR13064; TMEM9; 1.
Pfam; PF03128; CXCXC; 1.
Pfam; PF05434; TMEM9; 1.
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                                                                                                                                                                                                                                                                                                       Hypothetical protein; CHAIN 1 >18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GO; GO:0005770; C:late endosome; GO; GO:0005764; C:lysosome; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ensembl; SINFRUG00000152263; Fugu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PIR; T30536; T30536.
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    -!- SIMILARITY: Belongs to the TMEM9 family.

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NCE 182 AA; 20009 MW; 0869B43C89CB7475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AF026198; AAC15584.1; -;
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7 Putative protein
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Pred. No. 1.7e
0; Mismatches
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                                                                                                                                      Score 32;
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                                                                                                                                                                                                                                 /FTId=PRO 0000221039 Potential.
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                                                                                              Mismatches
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1.7e+02;
                                                                                                                                        DB 1;
                                                                                                               7e+02;
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                                                                                                                                   Length 187;
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RESULT 33 Q6CXD3_KLULA

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Q6ZRV1 HUD
ID Q6ZRV
AC Q6ZRV
DT 05-U
DT 05-U
DT 05-U
DT 05-U
DT 21-FF
DE CDNA
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Best Local S
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X PubMed=15229592; DOI=10.1038/nature02579;

X Dujon B., Sherman D., Fischer G., Durrens P., Casaregola S.,

Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,

A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,

A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,

Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,

A Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,

Bospons L., Fabre E., Faithead C., Ferry-Dumazet H., Groppi A.,

A Lespons L., Fabre E., Faithead C., Ferry-Dumazet H., Groppi A.,

A Lespons L., Fabre E., Faithead C., Ferry-Dumazet H., Groppi A.,

A Lespons L., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,

A Miraud J.-M., Nikolaki M., Oztas S., Ozier-Kalogeropoulos O.,

A Kerrest A., Koszul R., Wesolowski-Louvel M., Westhof E., Wirth B.,

A Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,

Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukhara M., Thierry A.,

Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,

Wincker P., Souciet J.-L.;

"Genome evolution in yeasts.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Complete
SEQUENCE
Ninomiya K., Wagatsuma M., Kanda K., Kondo H., Yokoi T., Kodaira Furuya T., Takahashi M., Kikkawa E., Omura Y., Abe K., Kamihara F. Katsuta N., Sato K., Tanikawa M., Yamazaki M., Sugiyama T., Irie Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J., Isono Y. Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H.,
                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004, integrated into UniProtKB/TrEMBL.
05-JUL-2004, sequence version 1.
21-FBB-2006, entry version 13.
CDNA FLJ46073 fis, clone TESTI2001364, highly similar lactate dehydrogenase A -like (LDHL).
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Sacci
Saccharomycetales; Saccharomycetaceae; Kluyveromyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-FEB-2006, entry version 12.
Similar to sp|P38869 Saccharomy
OrderedLocusNames=KLLA0A09295g;
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Q6CXD3;
                                                                                                                                             TISSUE=Testis;
                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
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Q6ZRV1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 430:35-44(2004).
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                                                                                                                                                                         NUCLEOTIDE SEQUENCE
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226 AA;
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WLLTAFS 72
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Euarchontoglires; Primates;
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Pred. No. 2.1e
2; Mismatches
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rimates; Catarrhini;
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InterPro; IPR011304; L_LDH_NAD.
InterPro; IPR011304; L_LDH_NAD.
InterPro; IPR001236; lact mal_DH.
InterPro; IPR000205; NAD_BS.
InterPro; IPR000594; Thif NAD_FAD_t
PANTHER; PTHR11540; ldh; I.
Pfam; PF002866; Ldh; I.C; 1.
Pfam; PF00056; Ldh; I.N; 1.
MUCLES (151 / OBRKOSH,
STRAIN=CDC 1551 / OBRKOSH,
MEDLINE=22206494; PubMed=12218036;
MEDLINE=22206494; PubMed=12218036;
DOI=10.1128/JB.184.19.5479-5490.2002;
DOI=10.1128/JB.184.19.5479-5490.2002;
                                                     NUCLEOTIDE
STRAIN=CDC
                                                                                                                                     Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C.M., Harris D.E., Gordon S.V., Eigimeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Davies R.M., Devlin K., Feltwell T., Gentles S., Hamlin Holroyd S., Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L.D., Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., Rutter S., Seeger K., Skelton S., Squares R., Rogers J., Rutter S., Seeger K., Skelton S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B.G.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSSP;
                                                                                                "Deciphering the biology of Mycobacterium complete genome sequence.";
Nature 393:537-544(1998).
                                                                                                                                                                                                                                                                                                            Mycobacterium tuberculosis NCBI_TaxID=1773;
                                                                                                                                                                                                                                                                                                                                                       Mycobacterium tuberculosis.
Bacteria; Actinobacteridae; Actinomycetales;
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                                                                                                                                                                                                                                                                                      NUCLEOTIDE
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243 AA; 26419 MW; 98D3B92406208F25 CRC64;
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                                                     SEQUENCE [LARGE 1551 / Oshkosh;
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                                                                                                                                                                                                                                                        PubMed=9634230; DOI=10.1038/31159;
                                                                                                                                                                                                                                                                                    [LARGE SCALE GENOMIC DNA].
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85.7%;
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                                                                     GENOMIC
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Attribution-NoDerivs License
              Carpenter
                                                                                                                             tuberculosis
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., Nagai K.,
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Complete
SEQUENCE
                                                                                                                                                                                      Garnier T., Eiglmeier K., Camus J.-C., Medina N., Mansoor H., Pryor M., Duthoy S., Grondin S., Lacroix C., Monsempe C., Simon S. Harris B., Atkin R., Doggett J., Mayes R., Keating L., Wheeler P.) Parkhill J., Barrell B.G., Cole S.T., Gordon S.V., Hewinson R.G., "The complete genome sequence of Mycobacterium bovis."; Proc. Natl. Acad. Sci. U.S.A. 100:7877-7882(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-2003, integrated into UniProtKB/TrEMBL.
01-OCT-2003, sequence version 1.
07-FEB-2006, entry version 11.
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GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0043236; F:ATPase activity, coupled
GO; GO:0006810; P:transport; IEA.
InterPro; IPR000412; ABC 2.
Pfam; PF01061; ABC2 membrane; 1.
Complete proteome.
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Gill J., Mikula
Fraser C.M.;
                                                                                                                                                                                                                                                                                                                                                        Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Corynebacterineae; Mycobacteriaceae; Mycobacterium; Mycobacterium tuberculosis complex.
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Hickey E.K., Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M.D.,
Salzberg S.L., Delcher A., Utcerback T.R., Weidman J.F., Khouri H.M.,
Gill J., Mikula A., Bishai W., Jacobs W.R. Jr., Venter J.C.,
                                                                                                                                              Copyrighted Distributed
                                                                                                                                                                                                                                                                                MEDLINE=22709107; PubMed=12788972; DOI=10.1073/pnas.1130426100;
                                                                                                                                                                                                                                                                                                                                                                                                                   Name=rfbD; OrderedLocusNames=Mb3812;
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                                                                                                                                                                                                                                                                                                                                                                                                     Mycobacterium bovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tuberculist; Rv3783;
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                          Pfam; PF01061; ABC2_membrane; 1.
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                                           GO:0006810; P:transport; IEA.
erPro; IPR000412; ABC_2.
                                                                    ; BXZ48347; CAD95998.1; -; Genomic_DNA.
GO:0016020; C:membranc; IEA.
GO:0005524; F:ATPase activity, coupled
GO:0042626; F:ATPase activity,
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                                                                                                                                             by the UniProt Consortium, see http://www.uniprot.org/terms under the Creative Commons Attribution-NoDerivs License
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 32336 MW;
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71.4%;
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Pred. No. 2.
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 D8793899C05A7EC3 CRC64;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AY042212; AAK91803.1; -; Genomic_DNA.EMBL; AJ313465; CAC86130.1; -; mRNA.Ensembl; ENSMUSG00000050276; Mus musculus.MGI; MGI:3033145; Mrgprg.
InterPro; IPR000276; GPCR_Rhodpsn.Pfam; PF00001; 7tm 1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyrighted by the UniProt Consortium, see nrcp://www.uniproc.org/consortium
Distributed under the Creative Commons Attribution NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The BWS region contains an evolutionary breakpoint regi
Submitted (JUL-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Engemann S., Stroedicke M., Meguro M., Oshimura M., Walter J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=21435808; PubMed=11551509; DOI Dong X., Han S.-K., Zylka M.J., Simon "A diverse family of GPCRs expressed i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria; Euarchontogl
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=Mrgprg; Synonyms=Ebrt2, Mrgg;
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -I- FUNCTION: Orphan receptor. May regulate nociceptor function development, including the sensation or modulation of pain.
-i- SUBCELULAR LOCATION: Membrane; multi-pass membrane protein.
-i- SIMILARITY; Belongs to the G-protein coupled receptor 1 fami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=C57BL/6;
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Pred. No.
   Cytoplasmic (Potential).
4 (Potential).
Extracellular (Potential)
5 (Potential).
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mon M.I., Anderson D.J.;
ed in specific subsets of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Receptor; Transducer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             see http://www.uniprot.org/terms
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AS Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Wakernan K.J., Malek J.A., Gunarathe P.H.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Kliausner R.D., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Rodrigues S., Sanchez A.,

Whiting M., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Hahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Hahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Balakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

Balakesley R.W., Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N.S., Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J. Schein J.E., Jones S.J.M., Marra M.A.;

Brock Matl N. Schein J. Schein J.E., Jones S.J.M., Marra M.A.;

Brock M. Schein J. Schein J. Schein J. S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=Mrgprg;
Mus musculus (Mouse).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Butheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;

Muroidea; Muridae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPO DOM
TRANSMEM
TOPO DOM
TRANSMEM
TOPO DOM
CONFLICT
EMBL; BC100302; AAI00303.1; -; mRNA.

MGI; MGI:3033145; Mrgprg.

GO; GO:0016021; C:integral to membrane; IEA.

GO; GO:0016020; C:membrane; IEA.

GO; GO:0004872; F:receptor activity; IEA.

GO; GO:0001584; F:rhodopsin-like receptor activity;

GO; GO:0007186; P:G-protein coupled receptor proteir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MAS-related GPR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q498A2 MOUSE
Q498A2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                       Distributed under the Creative Commons
                                                                                                                                                                                                                                                                                    Copyrighted by the UniProt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rissuE=Placenta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                           MGC Project; mitted (AUG-2005) to the EMBL/GenBank/DDBJ databases. SUBCELLULAR LOCATION: Membrane; multi-pass membrane
                                                                                                                                                                                                                                                                                                                                                    similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B
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71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     U.S.A.
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                                                                                                                                                                                                                                                                                       Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytoplasmic (Po
T -> I (in Ref.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytoplasmic (Potential).
6 (Potential).
Extracellular (Potential)
7 (Potential).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99:16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    82BD4BEA02E0864C CRC64;
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2.7e+02;
                                                                                                                                                                                                                                                    see http://www.uniprot.org/terms
Attribution-NoDerivs License
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         protein
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         signalin.
                                                                                                                                                                                                                                                                                                                                                                              protein
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RESULT 40
Q2TWR7 9DIPT
ID Q2TWR7;
ID Q2TWR7;
AC Q2TWR7;
DT 24-JAN-2006, s
DT 07-MAR-2006, s
DT 08-MAR-2006, s
DT 08-MAR-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q3M9S1 ANAVT
Q3M9S1;
25-OCT-2005,
25-OCT-2005,
07-FEB-2006,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US DOE Joint Genome Institute;
Copeland A., Lucas S., Lapidus A., Barry K., Detter J.C.,
Hammon N., Israni S., Pitluck S., Saunders E.H., Schmutz
Larimer F., Land M., Kyrpides N., Mavrommatis K., Richard
"Complete sequence of Anabaena variabilis ATCC 29413.";
Submitted (SEP-2005) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR000276; GPCR_Rhodpsn. Pfam; PF00001; 7tm_1; 1. PRINTS; PR00237; GPCRRHODOPSN.
                                                                                                                                                                                               24-JAN-2006, sequence version 07-MAR-2006, entry version NADH dehydrogenase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical
ORFNames=Ava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G-protein coupled receptor; Membrane; Receptor; Transducer;
                                                          Ephydroidea; Droso
NCBI_TaxID=311483;
                                                                                 Neoptera; Endopterygota; Dip
Ephydroidea; Drosophilidae;
                                                                                                                                                         Mitochondrion.
                                                                                                                                                                             Drosophila unimaculata.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical protein; SIGNAL 1 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; CP000117; ABA22265.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=ATCC 29413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anabaena varīabilis (strain ATCC 29413).
Bacteria; Cyanobacteria; Nostocales; Nos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
  NUCLEOTIDE SEQUENCE. Wang B.-C., Park J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=240292;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANAVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GO:0007165; P:signal transduction;
                                                                                                                                                                                                                                                                                                                                                                                                                    17
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                                                                                                                                                                                                                                                                                                                                                                                                                    WLLSSFS
                                                                                                                              Metazoa; Arthropoda;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               333 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   289 AA;
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2652;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry version 3.
                                                                                                                                                                                                                                                               integrated into UniProtKB/TrEMBL
                                                                                                                                                                                                                                          sequence version
                                                                                                                                                                                                                                                                                                            PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  84.2%;
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71.4%;
      Watabe
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                                                                                                       Diptera;

    Fragment).

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; 1F1E00BAF83097F3 CRC64;
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Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 32; DB 2;
Pred. No. 3.1e+02;
                                                                                                                                                                                                                                                                                                            PRT;
      . -A.,
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                                                                                                          Hexapoda; Insecta; Pterygota;
a; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
      Gao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nostocaceae; Anabaena.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    333
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      J.-J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
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RESULT 42
Q2TNS6_9DIPT
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Matches
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Best Local :
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IS2 9DIPT
Q2TNS2 9DIPT
Q2TNS2;
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24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fragment).
Drosophila medioconstricta.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Phylogenetic analysis of the Drosophila robusta and melanica species groups based on mitochondrial and nuclear DNA sequences."; Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol.
-!- SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner membrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mitochondrion; NAD; Oxi
                                                                                                                                                                                                                                                                                                                                                             EMBL; AY750079; AAX52010.1; -; Genomic_DNA.
EMBL; AY750074; AAX52005.1; -; Genomic_DNA.
GO; GO:0005739; C:mitochondrion; IEA.
GO; GO:0005739; Ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
Wang B.-C., Park J., Watabe H.-A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GO:0005739; C:mitochondrion; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              multi-pass membrane protein (By similarity).
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                                                                                                                                WMLSALS 191
                                                                                                                                                                               WMLSAFS 7
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341 AA;
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               PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                       39472 MW;
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85.7%;
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85.7%;
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Pred. No. 3.2e+02;
D; Mismatches 1
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               PRT;
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                                                                                                                                                                                                                                                                       Length 341;
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RESULT
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Best Local
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24-JAN-2006,
07-MAR-2006,
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T2 9DIPT
Q2TNT2 9DIPT
Q2TNT2;
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                                                                                                                                                                                             groups based on mitochondrial and nuclear DNA sequ
Submitted (SEP-2004) to the EMBL/GenBank/DDBJ data
-!- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+)
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Ephydroidea; Drosophilidae;
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24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fr
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Wang B.-C., Park J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitochondrion.
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                  GO; GO:0005739; C:mitochondrion; IE/
Mitochondrion; NAD; Oxidoreductase;
NON TER 341 341
                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/terms Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium, see http://www.uniprot.
Distributed under the Creative Commons Attribution-NoDerivs Li
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                                                                  EMBL; AY750069; AAX52000.1; -; Genomic_DNA.GO; GO:0005739; C:mitochondrion; IEA.
                                                                                                                                                                                                                                                                                                                                  Wang B.-C.,
Zhang Y.-P.;
                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
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    -:- SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner

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  SEQÜENCE
                                                                                                                                                                                                                                                                                                                 "Phylogenetic analysis of the Drosophila robusta and
                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=65966;
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                                                                                                                                                                                                                                                                                                                                                            Park J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     integrated into UniProtKB/TrEMBL.
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    39589
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  MW.
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Pred. No. 3.2e+02;
0; Mismatches 1
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  A9B3CC084A3B5B95 CRC64;
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                                                                                                                                                                                                                                                                                                                                                          Gao J.-J.,
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                                          Ubiquinone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  341 AA.
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                                                                                                                                                                                                                                                                                                                                                          Chen H.-W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             License
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,,
                                                                                                                                                                                                                          membrane;
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RESULT 45
Q2TNT5_9DIPT
ID Q2TNT5;
ID Q2TNT5;
DT 24-JAN-2006, s
DT 07-MAR-2006, s
DT 08-MAR-2006, s
DT 08-MAR-2006, s
DT 08-MAR-2006, s
DT 08-MAR-2006, s
DT 24-JAN-2006, s
DT 
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Best Local
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Best Local
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NUCLEOTIDE SEQUENCE.
Wang B.-C., Park J., Watabe H.
Zhang Y.-P.;
"Phylogenetic analysis of the
groups based on mitochondrial
                                                                                                                                                                                                                                                                                         24-JAN-2006, integrated into UniProtKB/
24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Phylogenetic analysis of the Drosophila robusta and melanica species groups based on mitochondrial and nuclear DNA sequences."; Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol.
-!- SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JAN-2006, integrated into uni
24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
                                                                                                                                                                      Neoptera; Endopterygota; Dip
Ephydroidea; Drosophilidae;
                                                                                                                                                                                                                                                                              Drosophila moriwakii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AY750068; AAX51999.1; -; Genomic_DNA.
GO; GO:0005739; C:mitochondrion; IEA.
Mitochondrion; NAD; Oxidoreductase; Ubiquinone.
NON_TER
341
341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Distributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Copyrighted by the UniProt Consortium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ephydroidea; Drosc
NCBI_TaxID=311477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila paramelanica.
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NADH dehydrogenase subunit 2 (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QZTNT3_9DIPT
QZTNT3;
                                                                                                                                                                                                                         Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wang B.-C., Park J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PDIPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                341 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84.2%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39575 MW;
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85.7%;
                                                                       Watabe H.-A.,
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                                                                                                                                                                                                                                                                                                                                                                             into UniProtKB/TrEMBL
                                                                                                                                                                                                  Diptera; Brachycera; Muscomorpha;
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Pred. No. 3.2e
0; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
     Drosophila robusta and melanica species and nuclear DNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4A1428DBA3BE5EBD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gao J.-J., Aotsuka
                                                                       Gao J.-J., Aotsuka T.,
                                                                                                                                                                                                                                                                                                                                                                                                                           341 AA.
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. 3.2e+02;
. 3.7 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .2e+02;
  DNA sequences.
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RESULT 47
Q2TNT7_DROMX
ID Q2TNT7;
AC Q2TNT7;
DT 24-UAN-2006, E
DT 24-UAN-2006, E
DT 07-MAR-2006, E

entry version 3.

integrated into UniProtKB/TrEMBL

PRELIMINARY;

PRT;

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RESULT 46
Q2TNT6 9D1
ID Q2TNT
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                                             Query Match
Best Local S
Matches 6
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Matches
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24-JAN-2006,
07-MAR-2006,
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GO; GO:0005739; C:mitochondrion; IEA.
Mitochondrion; NAD.
NON TER
341
341
SEQUENCE 341 AA; 39646 MW; 5D1F0BAE2786B4A5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AY750066; AAX51997.1; -; Genomic_DNA. GO; GO:0005739; C:mitochondrion; IEA. Mitochondrion; NAD; Oxidoreductase; Ubiquin NON_TER 341
                                                                                                                                                                                                                       "Phylogenetic analysis of the Drosophila robusta and melanica groups based on mitochondrial and nuclear DNA sequences."; Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databases.
-i- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol
                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Arthrop
Neoptera; Endopterygota; Di
Ephydroidea; Drosophilidae;
                                                                                                                                                                                                                                                                                                                                                                                         07-MAR-2006, entry version NADH dehydrogenase subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Copyrighted by the UniProt Consortium, see http://www.uniprot.org/to
                                                                                                                                                               Copyrighted by the UniProt Consortium, see http://www.uniprot.org/termsDistributed under the Creative Commons Attribution-NoDerivs License
                                                                                                                                                                                                                                                                                  Wang B.-C
                                                                                                                                                                                                                                                                                                                NCBI_TaxID=189258;
                                                                                                                                                                                                                                                                                                                                                                   Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                               Drosophila micromelanica.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Q2TNT6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQÜENCE
                                                                                                                                                                                                              -!- SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner
                                                                                                                                                                                                                                                                    Zhang Y.-P.;
                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              9DIPT
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                                                                                                                                                                                                multi-pass membrane protein (By similarity).
185
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                    ۲
                                            l Similarity
6; Conserv
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WMLSALS 191
                      WMLSAFS
                                                                                                                                                                                                                                                                                                                                          Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; 
3ndopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      341 AA;
                                                                                                                                                                                                                                                                                  Park J.,
                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 sequence version 1.
                                                                                                                                                                                                                                                                                                                                                                                                                           integrated into UniProtKB/TrEMBL
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                                                        84.2%;
85.7%;
                                                                                                                                                                                                                                                                                  Watabe H.-A., Gao J.-J.,
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                                                                                                                                                                                                                                                                                                                                                                                          (Fragment).
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Pred. No. 3
                                                         Score 32; DB 2;
Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         686D590E56FDE76A CRC64;
                                                                                           5D1F0BAE2786B4A5 CRC64;
                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  341
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                                                                                                                                                                                                                                                                                   Aotsuka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 341;
                                                                    Length 341;
                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                         ubiquinol.
                                                                                                                                                                                                                                                                                  T., Chen H.-W.,
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                                                                                                                                                                                                              membrane;
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                                                                                                                                                                                                                                                            species
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RESULT 48
Q2TNY9 9D1
ID Q2TNY1
AC Q2
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Query Match
Best Local S
Matches 6
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Phylogenetic analysis of the Drosophila robusta and melanica species groups based on mitochondrial and nuclear DNA sequences."; Submitted (SEP-2004) to the EMEL/GenHank/DDBJ databases -i- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-JAN-2006, integrated into Uni
24-JAN-2006, sequence version 1.
07-MAR-2006, entry version 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QZTNT9_9DIPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Phylogenetic analysis of the Drosophila robusta and melanica s groups based on mitochondrial and nuclear DNA sequences."; Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databbases.
-!- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol.
                                                                                                                                               NON TER
SEQUENCE
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Distributed
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NADH dehydrogenase subunit 2 (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota;
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                                                                                                                                                                                                                                                     EMBL; AY750062; AAX51993.1; -; Genomic_DNA.GO; GO:0005739; C:mitochondrion; IEA.
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Mitochondrion; NAD; Oxidoreductase; U
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                                                                                                                                                                                                                   Mitochondrion; NAD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhang Y.-P.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=65966;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (By similarity).
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Similarity 6; Conserv
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341 AA;
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                                                                                                                                                                                                                                                                                                                         by the UniProt Consortium, see http://www.uniprounder the Creative Commons Attribution-NoDerivs
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                                                                                                                                               39512 MW;
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                                     84.2%;
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Pred. No. 3.2e+02;
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                                     Score 32; DB 2;
Pred. No. 3.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                             http://www.uniprot.org/terms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aotsuka T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aotsuka T.,
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RESULT 50
Q2TNU1 9DIPT
ID Q2TNU1 9DIPT
AC Q2TNU1;
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Q2TNU0_9DIPT
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Best Local S
Matches 6
24-JAN-2006, integrated into UniProtKB/T
24-JAN-2006, sequence version 1.
27-JANR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fragment).
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Mitochondrion; NAD; Oxidoreductase; Ubiquinone
NON_TER 341 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    groups based on mitochondrial and nuclear DNA sequences.";
Submitted (SEP-2004) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: NADH + ubiquinone = NAD(+) + ubiquinol.
-!- SUBCELLULAR LOCATION: Mitochondrion; mitochondrial inner membrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-MAR-2006, entry version 3.
NADH dehydrogenase subunit 2 (Fragment)
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07-MAR-2006, entry version 3.
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                                                                                                                                              Wang B.-C.,
Zhang Y.-P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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Wang B.-C., Park J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ephydroidea; Drosophilidae; Drosophila
NCBI_TaxID=311474;
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                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                     Neoptera; Endopterygota; Diptera; Brachycera;
Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                          Park J.,
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Pred. No. 3.
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                                                                                                                                                                          Gao J.-J.,
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26 68.4 348 2 US-09-603-600-10 26 68.4 349 2 US-08-931-8974B-343 26 68.4 349 2 US-08-931-080-11 26 68.4 349 2 US-08-931-700-6 68.4 349 2 US-08-931-700-6 68.4 349 2 US-08-931-700-6 68.4 359 2 US-08-931-700-6 68.4 359 2 US-09-299-843-24 26 68.4 375 1 US-08-46-875-8 26 68.4 375 1 US-08-46-875-8 26 68.4 396 2 US-09-949-016-9465 20 US-09-949-016-9465 20 US-09-949-016-9465 20 US-09-605-703B-1020 26 68.4 396 2 US-09-605-703B-1020 26 68.4 422 2 US-09-604-855-114	26 68.4 325 2 US-08-158-735A-13 Sequence 1 68.4 329 2 US-09-603-208A-242 Sequence 2 668.4 331 2 US-09-603-208A-242 Sequence 2 668.4 345 2 US-09-603-208A-35 Sequence 2 26 68.4 345 2 US-09-193-681A-7546 Sequence 5 68.4 346 2 US-09-193-681A-7546 Sequence 6 68.4 346 2 US-09-193-33A-5 Sequence 2 668.4 346 2 US-09-993-33A-5 Sequence 3 68.4 346 2 US-09-993-33A-5 Sequence 3 68.4 346 2 US-09-993-112B-3 Sequence 3 68.4 346 2 US-09-09112B-3 Sequence 3 68.4 346 2 US-09-011-533-7 Sequence 3 68.4 346 2 US-09-011-533-7 Sequence 2 68.4 346 2 US-09-011-533-7 Sequence 3 68.4 346 2 US-09-011-533-7 Sequence 2 68.4 348 2 US-09-138-46 Sequence 2 68.4 348 2 US-09-138-46 Sequence 3 68.4 348 2 US-09-138-30 Sequence 3 68.4 348 2 US-09-138-10 Sequence 3 68.4 348 2 US-09-133-974B-342 Sequence 3 68.4 348 2 US-09-533-274B-342 Sequence

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US-09-949-016-10977
US-08-357-533A-10
US-08-459-009-11
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US-08-459-009-12
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Patent No. 6309877

GENERAL INFORMATION:

APPLICANT: Chau, Raymond M. W.

TITLE OF INVENTION: Isolation and Use of Mot
FILE REFERENCE: 12592-2

CURRENT APPLICATION NUMBER: US/08/928,862

CURRENT FILING DATE: 1997-09-12

NUMBER OF SEQ ID NOS: 5

SOPTWARE: PatentIN Ver. 2.0

SEQ ID NO 4

LENGTH: 33

TYPE: PRT

ORGANISM: Homo sapiens
RESULT 2
US-09-592-018-4
; Sequence 4, Application US/09592018
; Patent No. 6759389
; GENERAL INFORMATION:
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US-08-928-862-4
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Best Local S
Matches 7
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US-08-954-032-11
US-08-954-0368-19
US-08-954-0406-19
US-08-954-055-19
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Pred. No.
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US-07-757-022B-18
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US-09-989-481-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 09/633,447
PRIOR FILING DATE: 2000-08-07
PRIOR PELICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR FILING DATE: 1996-09-27
                                                                                                                                                                                                                                                                                 Best Loc
Matches
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 4
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APPLICANT: Chau, Raymond M.W.
TITLE OF INVENTION: Isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 38; Best Local Similarity 100.0%; Pred. No. Matches 7; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 4
LENGTH: 33
                                                                                                                       Sequence 18,
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Patent No. 6841531
                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 12592-3
CURRENT APPLICATION NUMBER: US/09/592,018
CURRENT FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1996-09-27
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chau, Ray TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2001-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 60/026792
                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 33
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                   APPLICANT:
                                   APPLICANT:
                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                              12 WMLSAFS 18
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                                                                                                       9, Application US/07757022B
6433142
   INVENTION:
                               Gesner, Thomas G. Clark, Stephen C. Turner, Katherine
                                                                                                                                                                                                                                                                                   Conservative
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Rodney M.
Megakaryocyte Stimulating Factors
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100.0%; Pred. No. 1.2;
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US-10-124-557-18
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                                                                                                                                                                                                                                           Sequence 18, Applicati
Patent No. 7030223
GENERAL INFORMATION:
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TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 18-JAN-19
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APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
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LENGTH: 46 amino acids
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TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US OF FILING DATE: 08-AUG-1989 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 87 Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: UPPLING DATE: 19910910
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
                                                                                                              NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM
                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                      1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AMINO ACID
GY: linear
                                             STREET: 87 CambridgePark Drive
CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                    WMLSPFS
               COUNTRY: U.S.A. ZIP: 02140
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                                                                                              ADDRESSEE: Genetics Institute,
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (617)876-1170
                                                                                                                                                                                                             Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-JAN-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       86.8%;
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Pred. No.
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RESULT 6
US-07-757-0228-72
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No. 6433142
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         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIFICATION = 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR DATE: 18-JAN-1991
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                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                  STREET: 87 Cambridge
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                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 46 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
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FILING DATE: 16-Apr-2002
CLASSIFICATION: -Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                  Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/07757022B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                    87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gesner, Thomas G. Clark, Stephen C.
                                                                                                                                                                                                                                                              U.S.A.
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                                                                                                                                                                                                                                                                                                                                       Genetics Institute, Inc.
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                                                                                                                                                                                                                                                                                                                                                                               143
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Pred. No. 15;
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US-10-124-557-72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 72
SEQUENCE CHARACTERISTICS:
LENGTH: 237 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 72, App.:.
Sequence 72, App.:.
No. 7030223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. 7030223
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION: NAME: Cserr, Luann
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02140
COMPUTER READABLE FORM:
                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                           CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  111 WMLSPFS 117
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                                                                APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 87 CambridgePark Drive CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
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(617)876-5851
'OR SEQ ID NO: 72:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER: US 07/390,901
08-AUG-1989
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Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hewick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clark, Stephen C.
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                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIPICATION: 530
PRIOR APPLICATION DATA: 300
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
PRIOR APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
PRIOR APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
PRIOR APPLICATION NUMBER: US 07/457,196
FILING DATE: 09-DEC-1989
PRIOR APPLICATION NUMBER: US 07/390,901
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
PRIOR APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                       TELEFAX: (617)876-585
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 72:
                                                                                                                                           NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                               SEQUENCE CHARACTERISTICS:
LENGTH: 372 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 86.8%;
Local Similarity 85.7%;
nes 6; Conservative
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TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 72:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 237 amino acids
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TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             111 WMLSPFS 117
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                           (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (617)876-5851
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 33; DB 3; Pred. No. 77;
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Query Match
Best Local Similarity
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Best Local Similarity
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US-10-124-557-64
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                                                                                                                                       ; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-124-557-64
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                                                                                                                                                                                                                                                TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARBE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA: Release #1.0,
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <undown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Megakaryocyte Stimulating Factors
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139 WMLSPFS 145
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                                 1 WMLSAFS 7
                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                        NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                               LENGTH: 372 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 87 CambridgePark Drive
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                                                                     Conservative
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Clark, Stephen C.
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                                                                                    86.8%;
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                                                                   Score 33; DB 3;
Pred. No. 1.2e+02;
0; Mismatches 1
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Pred. No. 1.2e+02;
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RESULT 11
US-10-124-557-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 68:
SEQUENCE CHARACTERISTICS:
                                                                                Sequence 68, Application US/10124557
Patent No. 7030223
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
PRIOR APPLICATION NUMBER: US 07/457,196
FILING DATE: 08-AUG-1989
PRIOR APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CREATY INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION: APPLICANT: Gesner
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                                                                                                                                                                                                                                                                                                                                                                      TYPE: AMINO A TOPOLOGY: lir MOLECULE TYPE:
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MEDIUM TYPE: Floppy of COMPUTER: IBM PC com
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CURRENT APPLICATION DATA:
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APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: b, Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                    APPLICANT:
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            Jacobs,
Hewick,
                                                 Turner, Katheri
Clark, Stephen
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85.7%;
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Rodney M. Thomas G.
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                                                                     Katherine
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Pred. No. 1.4e+02;
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TOPOLOGY: Intear

MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 68:
US-10-124-557-68
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Best Local Similarity
Watches 6; Conserve
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US-07-757-022B-66
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                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                           Sequence 66, Application US/07757022B
                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 68: SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                             APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakary
NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: Ceerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
                                                                                                                      STREET:
                                                                               COUNTRY:
                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             189 WMLSPFS 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 WMLSAFS 7
                                                                 02140
                                                                                                                        Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 422 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ELEPHONE: (617)876-1170
                                                                                                                                    87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                             Genetics Institute,
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Pred. No. 1.4e+02;
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OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,

Version #1.25

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Best Local Similarity
Watches 6; Conserva
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US-10-124-557-66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 66, Application US/10124557 Patent No. 7030223 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 423 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 423 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 29-DEC-1989

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 1:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: CSerr, Luann
NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX:
                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                          Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Turner, Katherine
Clark, Stephen C.
                               APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                190 WMLSPFS 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 WMLSAFS 7
                                                                                                                                                                                                                        STATE: Massachusetts
COUNTRY: U.S.A.
ZIP: 02140
                                                                                                                                                                                                                                                                                CITY: Cambridge
                                                                                                                                                                                                                                                                                                STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genetics Institute, Inc
APPLICATION NUMBER: US 07/643,502 FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                 Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19910910
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Pred. No. 1.4e+02;
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Best Local Similarity
Watches 6; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 54, Application US/07757022B Patent No. 6433142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 66:
SEQUENCE CHARACTERISTICS:
LENGTH: 423 amino acids
TYPE: amino acid
                                                                                                                                                                       CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/546,114
FILING DATE: 29-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gesner, Thomas G. APPLICANT: Clark, Stephen C. APPLICANT: Turner, Katherine APPLICANT: Hewick, Rodney M. TITLE OF INVENTION: Megakaryoc NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
            FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CBETT TARRET
                                                                      FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 66:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       190 WMLSPFS 196
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REFERENCE/DOCKET NUMBER: GI 5190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Megakaryocyte Stimulating Factors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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REFERENCE/DOCKET NUMBER: GI

5190

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; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: 54: US-10-124-557-54
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US-10-124-557-54
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Best Local Similarity
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                                                                                                                                      TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 54:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 463 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: GI 5190 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    230 WMLSPFS 236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Cambridge
STATE: Massachusetts
                                                        LENGTH: 463 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                         REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                  NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10124557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hewick, Rodney M. Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.A.
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Query Match
Best Local Similarity
Matches 6; Conserve
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US-07-757-022B-44
                              Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
                                                                                                                                                                                                   TELEFAX: (617)876-585
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT AFFILCATION NUMBER: US/07/75/, VALUE APPLICATION NUMBER: US/07/75/, VALUE FILING DATE: 19910910
CLASSIFICATION: 530
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 07/643,502
                                                                                                                                                                                                                                NAME: CSGTY, LUARN
REGISTATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 1270 amino acids
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                                  TOPOLOGY:
                                                                                                                                     LENGTH: 12,0
TVPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                230 WMLSPFS 236
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 1 WMLSAFS
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gesner, Thomas G.
Clark, Stephen C.
Turner, Katherine
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                                                                                                                                                                                                                  (617)876-5851
                                   Conservative
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                                                                                                                                   linear
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                                                 86.8%;
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Pred. No.
                                                 Score 33; DB 2;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                    5190
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                                   Mismatches
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1.5e+02;
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                                   1; Indels
                                                                  Length 1270;
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RESULT 18
US-07-757-022B-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-124-557-44
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                     Sequence 42, Application US/07757022B Patent No. 6433142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 44:
                          APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: CBERT, LUBRIN REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: OF COMPATION COMPATION

COMPUTER: OF COMPATION

MEDIUM

ADDITION

COMPUTER: OF COMPATION

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COMPUTER: OF COMPATION

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 WMLSAFS 7
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FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 1270 amino acids
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ilarity 85.7%;
Conservative
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Gesner, Thomas G.
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Pred. No. 4.2e+02;
0; Mismatches 1
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RESULT 19
US-10-124-557-42
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                                                                                                                                                                                                                                                                                                                     Sequence 42, Application Patent No. 7030223
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 85.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CGETT, LUADIN
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION NUMBER: US 07/546,114
PRIOR APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
PRIOR APPLICATION NUMBER: US 07/457,196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 1311 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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STREET: 87 Cambr
CITY: Cambridge
STATE: Massachus
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COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: AMINO ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE:
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ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
                                                                                                                                                                             Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                             APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WMLSAFS
                                                         CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Massachusetts
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (617)876-1170
                                                                                                                                                                                                                                                         Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/07/757,022B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 33; DB 2;
Pred. No. 4.4e+02
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RESULT 20
US-07-757-022B-142
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MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 42:
US-10-124-557-42
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAL PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-0AN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,512
FILING DATE: 18-0AN-1991
PRIOR APPLICATION NUMBER: US 07/546,114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 142, Application US/07757022B Patent No. 6433142
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Best Local Similarity 85.7%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 1311 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Gesner, Thomas G. APPLICANT: Clark, Stephen C. APPLICANT: Turner, Katherine APPLICANT: Hewick, Rodney M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                CITY:
                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1078 WMLSPFS 1084
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REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
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FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                  Massachusetts: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (617)876-5851
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Pred. No. 4.4e+02;
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RESULT 21
US-10-124-557-142
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 142, Applicat Patent No. 7030223 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION DATA: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CSEIT, LUAND:
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELEPHONE: (617)876-1170
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 142:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: AMINO ACID
TOPOLOGY: linear
MOLECULE TYPE: protein
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LENGTH: 1313 amino aci
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 29-JUN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
ATTORNBY/AGENT INFORMATION:
NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
                                                                                                                                                                          CLASSIFICATION: - Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, '
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1080 WMLSPFS 1086
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Megakaryocyte Stimulating Factors
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 WMLSAFS 7
                                                                           APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                            FILING DATE: 29-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Genetics Institute, STREET: 87 CambridgePark Drive
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Clark, Stephen C.
Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hewick, Rodney M. Gesner, Thomas G.
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Pred. No. 4.4e+02;
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                                                                                                                                                                                                                                                                                                                                        Version
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; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: 142: US-10-124-557-142
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US-07-757-022B-50
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                                                                                                                                     FILING DATE: 19910910

PILING DATE: 19910910

PRIOR APPLICATION 530

PRIOR APPLICATION UNMBER: US 07/643,502

FILING DATE: 18-JAN-1991

PRIOR APPLICATION NUMBER: US 07/546,114

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

PRIOR APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

PRIOR APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

PRIOR APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

PRIOR APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:

NAME: CSert, Luann

REGISTRATION NUMBER: G1 5190

TELEPHONE: (617)876-1170

TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 50, Application US/07757022B Patent No. 6433142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                    TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 142:
SEQUENCE CHARACTERISTICS:
LENGTH: 1313 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1314 amino aci-
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gesner, Thomas G. APPLICANT: Clark, Stephen C. APPLICANT: Turner, Katherine APPLICANT: Hewick, Rodney M.
TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITILE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B6.8%;
Local Similarity 85.7%;
les 6; Conservarium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Genetics Institute, Inc.
87 CambridgePark Drive
                                                             314 amino acids
                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (617)876-1170
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Pred. No. 4.4e+02;
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                                                                                                                                            ; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-124-557-50
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Best Local Similarity
Thehes 6; Conserve
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US-10-124-557-50
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GENERAL INFORMATION:
                                                                        Matches
                                                                                          Query Match
Best Local (
                                                                                                                                                                                                                                                                              TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                  PRICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
AFTORNEY/AGENT INFORMATION:
NAME: CBEET, LUANITION:
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION TELEPHONE; (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1081 WMLSPFS 1087
1081 WMLSPFS 1087
                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Megakaryocyte Stimulating Factors
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                                     1 WMLSAFS 7
                                                                      Similarity 6; Conserv
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STATE: Massachusetts
                                                                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                          LENGTH: 1314 amino acids
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Pred. No. 4.4e+02;
                                                                        Score 33; DB 3;
Pred. No. 4.4e+02;
0; Mismatches 1
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                                                                                                           Length 1314;
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                                                                        Gaps
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                                                                                                                                                                                    RESULT 25
US-07-757-022B-60
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                                                                                                                           Sequence 60, Application US/07757022B Patent No. 6433142 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-70AN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
PRIOR APPLICATION UNBER: US 07/457,196
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Patent No. 6433
                                                                                                                                                                                                                                                                                                                                                    Query Match 86.8%;
Best Local Similarity 85.7%;
Matches 6; Conservative
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APPLICANT: Gesner, Thomas G. APPLICANT: Clark, Stephen C. APPLICANT: Turner, Katherine APPLICANT: Hewick, Rodney M. TITLE OF INVENTION: Megakaryoc NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (617)876-585: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390
PILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5
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ADDRESSEE: Genetics Institute, Inc.
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STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 1320 LENGTH: 1320 ACID
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    Megakaryocyte Stimulating Factors
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Pred. No. 4.4e+02;
0; Mismatches 1;
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TYPE: AMINO ACID
TOPOLOGY: linear
HOLECULE TYPE: protein
US-07-757-022B-60
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US-10-164-595-58
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                                                                           GRGANISM: Homo sapiens US-10-164-595-58
                                                                                                                                                             Sequence 58, Application US/10164595
Patent No. 6657054
GENERAL INFORMATION:
APPLICANT: Origene Technologies, Inc
TITLE OF INVENTION: Regulated Angiogenesis Genes and Polypeptides
FILLE REFERENCE: 1U 103 R1
CURRENT FILING DATE: 2002-06-10
NUMBER OF SEQ ID NOS: 80
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 6; Conserv
 Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                               SEQ ID NO 58
LENGTH: 1320
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APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1320 amino aci
TYPE: AMINO ACID
                                                                                                               TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Cambridge
STATE: Massachusetts
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87 CambridgePark Drive
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     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    86.8%;
                  86.8%;
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   Score 33; DB Pred. No. 4.4e 0; Mismatches
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Pred. No. 4.4e+02;
Db _,
. 4.4e+02;
. ~ 1;
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                                     Length 1320;
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       Indels
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       0
   Gaps
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 46:
US-10-124-557-46
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US-10-124-557-46
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                                                  RESULT 28
US-10-124-557-60
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Sequence 60, Application US/10124557 Patent No. 7030223 GENERAL INFORMATION:
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                                                                                                                                                                                                Matches
                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617)876-58: INFORMATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compartible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: GI 5190 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                          1087 WMLSPFS 1093
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                          1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                             LENGTH: 1320 amino acids
                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hewick, Rodney M. Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EE: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                     (617)876-1170
                                                                                                                                                                                                                 86.8%;
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                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                               Score 33; DB 3;
Pred. No. 4.4e+02;
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RESULT 29
US-07-757-022B-48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                 GENERAL INFORMATION:
APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPAX: (617) 976-5851
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
LENGTH: 1320 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity tes 6; Conserv
                 STREET: 87 CambridgePark Drive CITY: Cambridge
                                                       ADDRESSEE:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                              1087 WMLSPFS 1093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Cserr, Luann
NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                         Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 4.4e+02;
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COUNTRY: U.S.A.

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RESULT 30
US-10-124-557-48
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Best Local Similarity 85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 29-DEC-1989
PRIOR APPLICATION NUMBER: US 07/390
APPLICATION NUMBER: US 07/390
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CSerr, Luarn
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
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                      ZIP: 02140
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1121 WMLSPFS 1127
                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6; Conservative
                                                                                                                                                                       STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                 STREET: 87 CambridgePark Drive CITY: Cambridge
APPLICATION NUMBER: US/10/124,557
                                                                                                                                                                                                                                                           ADDRESSEE: Genetics Institute, Inc
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                                                                                                                                                                                                                                                                                                                                          Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 4.5e+02;
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RESULT 31
US-07-757-022B-40
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Best Local Similarity
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                            CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
110 07/457 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gesner
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                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
APPLICATION NUMBER: US 0 FILING DATE: 29-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 87 Cambı
CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 48:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1121 WMLSPFS 1127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 16-Apr-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/546,114 FILING DATE: 29-JUN-1990
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                                                                                                                                                                                                                                                                                                                                                                                                Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                     87 CambridgePark Drive
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Megakaryocyte Stimulating Factors
                      US 07/457,196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 33; DB 3;
Pred. No. 4.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 1354;
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RESULT 32
US-10-124-557-40
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 40, Applicati
Patent No. 7030223
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 40:
INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
LENGTH: 1361 amino ac
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ATTORNEY/AGENT INFORMATION:
NAME: Cserr, Luann
                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 1361 amino acids TYPE: AMINO ACID TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/390,901
                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1128 WMLSPFS 1134
                                                         NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 87 Cambı
CITY: Cambridge
                                            TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genetics Institute, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86.8%;
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Pred. No. 4.5e+02;
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TOPOLOGY: linear;
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
US-10-124-557-40
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US-07-757-022B-52
                                                                        US-07-757-022B-52
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Patent No. 6433142
                                                                                                                       TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 52:
SEQUIENCE CHARACTERISTICS:
LENGTH: 1363 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local :
                                   Query Match
                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CEETT, LUARIN
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Gesner
APPLICANT: Clark,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 19910910
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UNBER: US 07/643,502
FILING DATE: 18-JAN-1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/757,022B
FILING DATE: 19910910
                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Turner, Katherine APPLICANT: Hewick, Rodney M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 86.8%;
Local Similarity 85.7%;
tes 6; Conservative
Local Similarity 85.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/07757022B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            87 CambridgePark Drive
                                                                                                                                                                            (617)876-5851
(617) 876-5851
TD NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
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Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genetics Institute, Inc.
                                                                                        protein
86.8%; Score 33; DB
85.7%; Pred. No. 4.50
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,</u>
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                                                                                                                                                                                                                                                            5190
               4.5e+02;
                                     DB 2;
                                 Length 1363;
 Indels
 0;
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WMLSAFS

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RESULT 34
US-10-124-557-52
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                                                                                RESULT 35
US-07-757-022B-2
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                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: linear ; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-124-557-52
Sequence 2, Application US/07757022B
Patent No. 6433142
GEMERAL INFORMATION:
APPLICANT: Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 52, Applicati Patent No. 7030223 GENERAL INFORMATION:
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 52: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                              B6.8%;
Local Similarity 85.7%;
hes 6; Conservarium
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION IMFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1130
                                                                                                                                                            1130 WMLSPFS 1136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Turner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                    1 WMLSAFS 7
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STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 1363 amino acids
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Hewick, Rodney M.
Gesner, Thomas G.
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                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                             Score 33; DB 3;
Pred. No. 4.5e+02;
                                                                                                                                                                                                                                                                                                                                              52:
                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                Length 1363;
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Query Match
Best Local Similarity 8
""" hes 6; Conservat!
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US-07-757-022B-62
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                                                                                                                                                                                                                   Sequence 62, Application US/07757022B Patent No. 6433142
                                                                                                                        GENERAL INFORMATION:
APPLICANT: Gesner, Thomas G.
APPLICANT: Clark, Stephen C.
APPLICANT: Turner, Katherine
APPLICANT: Hewick, Rodney M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617)876-5851 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Hewick, Rodney M.
APPLICANT: Hewick, Rodney M.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN 1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 29-DEC-
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                   ADDRESSEE: Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 1404 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 18-JAN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                       TITLE OF INVENTION:
NUMBER OF SEQUENCES:
STREET: 8, STREET: 8, Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 199109 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 87 Cambi
CITY: Cambridge
STATE: Massachus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 02140
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)GY: linear
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                                87 CambridgePark Drive
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                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Katherine
Rodney M.
                                                                                       Megakaryocyte Stimulating Factors: 143
                                                                                                                                                                                                                                                                                                                                                                                                                           86.8%;
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Pred. No. 4.7e+02;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1404;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy

Floppy disk

ZIP: 02140

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                                                                                                                                       ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-164-595-78
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US-10-164-595-78
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APPLICANT: OriGene Technologies, Inc
TITLE OF INVENTION: Regulated Angiogenesis Genes and Polypeptides
FILE REFERENCE: 1U 103 R1
CURRENT APPLICATION NUMBER: US/10/164,595
CURRENT FILING DATE: 2002-06-10
NUMBER OF SEQ ID NOS: 80
                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1 SEQ ID NO 78
                                                                                                                                                                                                                                                                                                                                                                        Sequence 78, Application US/10164595
Patent No. 6657054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 86.8%;
Best Local Similarity 85.7%;
                                                                   Matches
                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CBerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 62:
                                                                                                                                                                                           LENGTH: 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/390,901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19910910
CLASSIFICATION: 530
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APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
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1171 WMLSPFS 1177
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                               1 WMLSAFS 7
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AMINO ACID
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                                                                   Conservative
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                                                                                  86.8%;
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                                                               Score 33; DB 2; Length 1404; Pred. No. 4.7e+02; 0; Mismatches 1; Indels
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GENERAL INFORMATION:
APPLICANT: Gregory D. Jay
TITLE OF INVENTION: tribonectins
FILE REFERENCE: 21486-026cip
CURRENT APPLICATION NUMBER: US/09/556,246
CURRENT FILING DATE: 2000-04-24
PRIOR APPLICATION NUMBER: USSN 09/298/970
PRIOR FILING DATE: 199-04-23
NUMBER OF SEQ ID NOS: 34
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jay, Gregory D.
TITLE OF INVENTION: TRIBONECTINS
CURRENT APPLICATION NUMBER: US/09/298,970A
CURRENT FILING DATE: 2001-06-19
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
FENCTH. 1404
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT ; ORGANISM: Homo sapiens US-09-897-188-1
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Best Local Similarity 85..
Whiches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Jay, Gregory D.
APPLICANT: Jay, Gregory D.
TITLE OF INVENTION: Tribonectin Polypeptides and Uses Thereof
FILE REFERENCE: 21486-026 CIP2
CURRENT APPLICATION NUMBER: US/09/897,188
CURRENT FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: 09/298,970
PRIOR APPLICATION NUMBER: 09/256,246
PRIOR APPLICATION NUMBER: 09/556,246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09298970A Patent No. 6743774 GENERAL INFORMATION:
                                                                                                                                                                              Sequence 1, Application US/09556246 Patent No. 7001881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09897188 Patent No. 6960562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2000-04-24
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
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Pred. No. 4.7e+02;
0; Mismatches 1
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Pred. No.
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US-10-124-557-2
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; ORGANISM: Homo sapiens
US-09-556-246-1
                                                                                                             US-10-124-557-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/10124557 Patent No. 7030223 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 86.8%;
Best Local Similarity 85.7%;
Matches 6; Conservative
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LENGTH: 1404
                                       Matches
                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1171 WMLSPFS 1177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02140
COMPUTER READABLE FORM:
                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID
                                                                                                                                                                                                                                                                                      NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Turner,
                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                     6
   WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Genetics Institute, Inc. STREET: 87 CambridgePark Drive
                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Massachusetts
                                                                                                                                                                                                 LENGTH: 1404 amino acids
                                                                                                                                                                                                                                                      TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                         TELEPHONE:
                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Clark, Stephen C.
                                                                                                                                                                                                                                                                         (617)876-1170
                                                    86.8%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Katherine
                                     0;
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Pred. No. 4.7e+02;
                                   Score 33; DB 3;
Pred. No. 4.7e+02;
0; Mismatches 1
                                                                                                                              NO:
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                                     1; Indels
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; TOPOLOGY: linear; MOLECULE TYPE: protein; SEQUENCE DESCRIPTION: SEQ ID NO: 62: US-10-124-557-62
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US-10-124-557-62
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                                                                                                RESULT 43
US-09-949-016-10827
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Sequence 10827, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: FOLYMORPHISMS IN KNOWN GENES ASSOCIATED
                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 7030223
GENERAL INFORMATION:
APPLICANT: Turn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 62,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: «Unknown»

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-UN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/124,557

FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1171 WMLSPFS 1177
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                                                                                                                                                                                                                        1 WMLSAFS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 1404 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10124557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Turner,
                                                                                                                                                                                                                                                                               86.8%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Luann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Katherine
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                                                                                                                                                                                                                                                                               Pred. No. 4.
                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5190
                                                                                                                                                                                                                                                                               DB 3;
4.7e+02;
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                                                                                                                                                                                                                                                                                                  Length 1404;
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                                                                                                                                                                                                                                                               Gaps
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US-09-107-532A-4889
US-09-107-532A-4889;
Sequence 4889, Application US/09107532A

Patent No. 6583275

Patent INFORMATION:

APPLICANT: Lynn A Doucette-Stamm and David Bush

APPLICANT: Lynn A DOUCETTO ACID AMINO ACID SEQUENCES RELATING TO

ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Human US-09-949-016-10827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10827
LENGTH: 1411
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 207012
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CURRENT FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                              INFORMATION FOR SEQ ID NO: 4889: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: J14 /2 , 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: ASCII
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1178 WMLSPFS 1184
                                    MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                        TYPE: amino acids
                                                                                                                                                                                                                              TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02354
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                    ORGANISM: Enterococcus
                                                                                                   TOPOLOGY: linear
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                                                                                                                                                                                                              EFAX:
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                                                                                                                                                                                                              (781)893-8277
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Pred. No. 4.7e+02;
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APPLICANT: Elsenberg, David
APPLICANT: Rotstein, Sergio H.
APPLICANT: Marcotte, Edward M.
APPLICANT: Marcotte, Edward M.
TITLE OF INVENTION: DETERMINING THE FUNCTIONS ANI
TITLE OF INVENTION: INTERACTIONS OF PROTEINS BY
FILE REFERENCE: 07419-032001
CURRENT APPLICATION NUMBER: US/09/712,363
CURRENT FILING DATE: 2000-01-13
PRIOR APPLICATION NUMBER: PCT/US00/02246
PRIOR FILING DATE: 2000-01-26
PRIOR FILING DATE: 2000-01-26
PRIOR APPLICATION NUMBER: 60/179,531
PRIOR APPLICATION NUMBER: 60/17,844
PRIOR FILING DATE: 1909-01-29
PRIOR APPLICATION NUMBER: 60/117,844
PRIOR APPLICATION NUMBER: 60/117,844
PRIOR APPLICATION NUMBER: 60/117,849
PRIOR APPLICATION NUMBER: 60/118,206,
PRIOR APPLICATION NUMBER: 60/118,206,
PRIOR FILING DATE: 1999-01-29
PRIOR FILING DATE: 1999-01-29
PRIOR FILING DATE: 1999-01-29
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US-09-252-991A-22359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-252-991A-22359
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SEQ ID NO 22359
LENGTH: 262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22359, Application US/09252991A Patent No. 6551795
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/252,991A CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/074,788 PRIOR FILING DATE: 1998-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERADERITY CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: UNSURE LOCATION: (182)
OTHER INFORMATION: Identity of amino acid at the above locations are unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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LOCATION: (B) LOCATION 1...103
SEQUENCE DESCRIPTION: SEQ ID NO: 4889:
                                   APPLICATION NUMBER: 60/126,593
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            234 WMLSAIS 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS 107196.136
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60/134,093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 32;
Pred. No.
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Pred. No.
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51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 262
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Sequence 99, Application US/09586106D

Patent No. 6720479

GENERAL INFORMATION:

APPLICANT: Wright, David A.

APPLICANT: Wyotas, Daniel F.

TITLE OF INVENTION: PLANT RETROCLEMENTS AND METHODS RELATED THERETO
FILE REFERENCE: P-1065A

CURRENT APPLICATION NUMBER: US/09/586,106D

CURRENT FILING DATE: 2003-02-07

PRIOR APPLICATION NUMBER: 60/087,125

PRIOR APPLICATION NUMBER: 69/087,125

PRIOR APPLICATION NUMBER: 09/322,478

PRIOR FILING DATE: 1998-05-29

PRIOR FILING DATE: 1999-05-28

NUMBER: OF SEQ ID NOS: 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 47
US-09-489-039A-7403
; Sequence 7403, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Mycobacterium tuberculosis US-09-712-363-283
                                                                                                                                                                                                                                                                                                                                                                                RESULT 48
US-09-586-106D-99
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LENGTH: 437
; TYPE: PRT
; ORGANIZM: Klebsiella pneumoniae
US-09-489-039A-7403
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SEQ ID NO 99
LENGTH: 229
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PUBLIMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 7403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 292
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 283
LENGTH: 280
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/165,124
PRIOR FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: 60/165,086
PRIOR FILING DATE: 1999-11-12
                                                             SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Y Match 84.2%;
Local Similarity 85.7%;
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APPLICATION NUMBER: 60/
FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      391 WSLSAFS 397
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60/134,092
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Pred. No. 1.4e+02;
2; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 32; DB 2;
Pred. No. 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                              Sequence 3700, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 190
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 99
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APPLICANT: Voytas, Daniel F.
APPLICANT: Voytas, Daniel F.
INTER OF INVENTION: PLANT RETROELEMENTS AND METHODS RELATED THERETO
FILE REFERENCE: P-1065A
CURRENT APPLICATION NUMBER: US/10/799,870
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: US/09/586,106
PRIOR FILING DATE: 2003-02-07
PRIOR APPLICATION NUMBER: 60/087,125
PRIOR PILING DATE: 1998-05-29
PRIOR PILING DATE: 1998-05-29
PRIOR PILING DATE: 1998-05-28
PRIOR FILING DATE: 1999-05-28
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TYPE: PRT
COMPUTER: PC
OPERATING SYSTEM: <Unknown>
OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
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MEDIUM TYPE: CD/ROM ISO9660
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1; Mismatches 1
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Pred. No. 1.8e+02;
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APPLICATION NUMBER: 60/051571

FILING DATE: July 2, 1997

ATTORNEY/ACENT INFORMATION:

NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489

REFERENCE/DOCKT NUMBER: GTC-012

TELECHONICATION INFORMATION:

TELEPAX: (781)893-8007

TELEPAX: (781)893-8277

INFORMATION FOR SEQ ID NO: 37000:

SEQUENCE CHARACTERISTICS:

LENGTH: 119 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: protein
HYDOTICIAL: YES
ORIGINAL SOURCE:
ORIGINAL SOURCE:
ORIGINAL SOURCE:
ORIGINAL SOURCE:
ORIGINAL SOURCE:
ORIGINAL SOURCE:
INFORMATION: (8) LOCATION 1...119

SEQUENCE DESCRIPTION: SEQ ID NO: 3700:
US-09-107-532A-3700

Query Match
PEATURE:
NAME: Arinierly 71.4%; Pred. No. 1.4e+02;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps
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Search completed: August 29, 2006, 06:17:48

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US-10-541-343-3

; Sequence 3, Application US/10541343

; Publication No. US20060052299A1

; GENERAL INFORMATION:

APPLICANT: Chau, Raymond Ming Wah
APPLICANT: KO, Pui.Yuk Dorochy

TITLE OF INVENTION: MNTF Peptides and Compositi
FILE REFERENCE: 2001-103US

CURRENT FILING DATE: 2005-07-05

PRIOR APPLICATION NUMBER: DCT/US2004/001468

PRIOR FILING DATE: 2004-01-21

PRIOR APPLICATION NUMBER: 60/441,722

PRIOR APPLICATION NUMBER: 60/441,722

PRIOR APPLICATION NUMBER: 00/441,722

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PRIOR APPLICATION NUMBER: 60/441,722
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Best Local
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                WMLSAFS
                                                                    100.0%; Score 38; ilarity 100.0%; Pred. No. Conservative 0; Mismatch
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US-10-325-175-307311

US-10-335-977-9307

US-09-864-761-45039

US-10-325-115-266020

US-10-425-115-266021

US-10-425-115-266021

US-10-425-115-266021

US-10-424-599-218746

US-10-02-344A-168

US-10-02-344A-168

US-10-02-344A-168

US-10-02-344A-168

US-10-02-344A-168

US-10-02-344A-168

US-10-02-34479-24

US-10-02-386-28417

US-09-833-245-1395

US-11-264-096-1396

US-11-264-096-1396

US-11-264-096-1396

US-11-424-599-23057

US-10-424-599-23057

US-10-424-599-230586

US-11-424-599-230586

US-11-425-115-304184

US-10-425-115-39447

US-10-425-115-39456

US-10-425-115-39469

US-10-425-115-39469

US-10-425-115-39457

US-10-425-115-39469
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                                                                          Mismatches
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                                                                                                     DB 5;
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                                                                                                Length
                                                                        Indels
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9307, Ap
9308, Ap
29471,
266020,
266021,
2449255,
168, App
1395, Ap
1395, Ap
1395, Ap
1395, Ap
1395, Ap
1395, Ap
23057, Ap
23057, Ap
230586, Ap
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US-10-541-343-6

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RESULT 4
US-10-858-545-4
; Sequence 4, Application US/10858545
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; OTHER INFORMATION: Synthetic peptide
US-10-541-343-6
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US-09-989-481-4
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CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: US 09/633,447
PRIOR EPILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR FILING DATE: 1996-09-27
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SEQ ID NO 4
LENGTH: 33
TYPE: PRT
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Publication No. US20060052299A1
GENERAL INFORMATION:
GENE
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Best Local Similarity
Matches 7; Conserv
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LENGTH: 13
TYPE: PRT
ORGANISM: Artificial
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Best Local
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CURRENT FILING DATE: 2005-07-05
PRIOR APPLICATION NUMBER: PCT/US2004/001468
PRIOR FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: 60/441,722
PRIOR PILING DATE: 2003-01-21
NUMBER OF SEQ ID NOS: 7
NUMBER OF SEQ ID NOS: 7
NUMBER OF SEQ ID NOS: 7
NUMBER: PatentIn version 3.3
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
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100.0%; Pred. No. 6.8;
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US-10-858-543-4

Sequence 4, Application US/10858543 Publication No. US20040266691A1 GENERAL INFORMATION:

Raymond M.W.

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; ORGANISM: Homo Sapiens
US-10-858-286-4
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CURRENT FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/09/592,018
PRIOR FILLING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR APPLICATION NUMBER: US 60/026792
                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.0
SEQ ID NO 4
LENGTH: 33
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Publication No. US20040242486A1
GENERAL INFORMATION:
APPLICANT: Chau, Raymond M.W.
TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-3
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                                                                     Matches
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PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
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PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
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CURRENT FILING DATE: 2004-06-01
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-3
                                                                                                                                                                                                                                                             PRIOR FILING DATE:
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NUMBER OF SEQ ID NOS: 5
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TYPE: PRT
ORGANISM: Homo Sapiens
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                    1 WMLSAFS 7
                                                                  Similarity 7; Conserv
 WMLSAFS 18
                                                                   Conservative
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                                                                                    Score 38;
Pred. No.
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                                                                     Mismatches
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CURRENT FILING DATE: 2004-06-01
PRIOR APPLICATION NUMBER: US/09/592,018
PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-01-15
PRIOR FILING DATE: 1996-09-27
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US-10-858-543-4
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PRIOR FILING DATE: 2000-06-12
PRIOR APPLICATION NUMBER: US 08/9928862
PRIOR FILING DATE: 1997-09-12
PRIOR APPLICATION NUMBER: US 08/751225
PRIOR FILING DATE: 1996-11-15
PRIOR APPLICATION NUMBER: US 60/026792
PRIOR FILING DATE: 1996-09-27
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Best Local S
Matches 7
                                                                                     Sequence 1, Application US/10541343
Publication No. US20060052299A1
GENERAL INFORMATION:
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SEQ ID NO 4
LENGTH: 33
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SEQ ID NO 4
LENCTH: 33
TYPE: PRT
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Publication No. US20060025565A1
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APPLICANT: Chau, Raymond Ming Wah
APPLICANT: Ko, Pui-Yuk Dorothy
TITLE OF INVENTION: MYTE Peptides and Compositions and Methods of Use
FILE REFERENCE: 2001-103US
                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-858-144-4
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TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
FILE REFERENCE: 12592-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                           y Match 100.0%; Score 38; DB Local Similarity 100.0%; Pred. No. 6.8; hes 7; Conservative 0; Mismatches
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                                                                                                                                                                          US-11-097-143-16893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR FILING DATE: 1999-10-19
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-28
PRIOR PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-12
PRIOR FILING DATE: 1999-11-12
PRIOR FILING DATE: 1999-11-12
                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 43008
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16893, Application US/11097143
Publication No. US200502085881
GENERAL INFORMATION:
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PRIOR FILING DATE: 2003-01-21
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.3
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 38; Best Local Similarity 100.0%; Pred. No. Matches 7; Conservative 0; Mismatch
                                                                                    Matches
                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Venter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS
TITLE OF INVENTION: ARRAYS, FOR DETECTING
TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
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CURRENT FILING DATE: 2005-07-05
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TYPE: PRT
ORGANISM: Artificial
FEATURE:
OTHER INFORMATION: Synthetic peptide
                                                                                                                                                                                               ORGANISM: DROSOPHILA
                                                                                                                                                                                                                                        LENGTH:
  966
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                                           1 WMLSAFS 7
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WMVSAFS
                                                                                    Conservative
  972
                                                                                                      Score 35; DB 6;
Pred. No. 8.4e+02;
                                                                                      Mismatches
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EXPRESSION OF 10,000 OR MORE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 5;
                                                                                                                             Length 1760;
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RESULT 10 US-10-424-599-187845

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RESULT 11
US-10-124-557-18
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US-10-424-599-187845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5323)B CURRENT APPLICATION NUMBER: US/10/424,599 CURRENT FILING DATE: 2003-04-28 NUMBER OF SEQ ID NOS: 285684 SEQ ID NO 187845 LENGTH: 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18, Application US/10124557 Publication No. US20020137894A1 GENERAL INFORMATION:
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                  APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
APPLICATION UNMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
REFERENCE/DOCKET NUMBER: GI 5190 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Genetics Institute, STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/643,502 FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Cambridge
                                           REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
                                                                     NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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Hewick, Rodney M.
Gesner, Thomas G.
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                                                                     Luann
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US-11-169-232-18
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Publication No. US2006
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 46 amino acids
                                                                                           INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 46 amino aci
                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/10/124,557

FILING DATE: 16-Apr-2002

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-UN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/457,196
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hes 6; Conserv
                                                                                                                                                                 REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION TELEPHONE; (617)876-1170
                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/390,901 FILING DATE: 08-AUG-1989 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                 MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
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Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                       TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                         NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 28-Jun-2005 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A. ZIP: 02140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Genetics Institute, STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                   TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/11169232
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Gesner, Thomas G.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                         GI 5190
                     18:
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RESULT 13
US-09-925-301-1452
; Sequence 1452, Application US/09925301
; Patent No. US20020052308A1
; GENERAL INFORMATION:
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; Sequence 72, Application US/10124557
; Publication No. US20020137894A1
; GENERAL INFORMATION:
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Rosen et al. TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE
LOCATION: (6)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 61
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 37 WMLSPFS 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ហ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6; Conservative
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                                                                                                                                                                             CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                              ZIP: 02140
                                                                                                                                                                                                                                    STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                      ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                           Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 33; DB 3;
Pred. No. 1e+02;
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                                                                        Version
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Query Match
Best Local Similarity
Matches 6; Conserva
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MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 72:
US-10-124-557-72
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US-11-169-232-72
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Publication No. US20060025570A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 72: SEQUENCE CHARACTERISTICS:
                                                                                                                          CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
CILOCATION NUMBER: US/11/169,232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/390,901 FILING DATE: 08-AUG-1989 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          111 WMLSPFS 117
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                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UNN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 237 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hewick, Rodney M. Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86.8%;
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Pred. No. 3.3e+02;
0; Mismatches 1
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Publication No. US20020137894A1
GENERAL INFORMATION:
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
                                            INFORMATION FOR SEQ ID NO: 64:
                                                                                                                                                                            APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 72:
                                                        TELEPAX: (617)876-5851
TELECOMMUNICATION INFORMATION:
TELEPAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Turner, Katherine Clark, Stephen C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6,
                                                                                                                                        NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 237 amino acids
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  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EB: Genetics Institute, Inc.
87 CambridgePark Drive
372 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      86.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 33; DB 6; Length 237; Pred. No. 3.3e+02;
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                                                               US-11-169-232-64
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                               Query Match
 Matches
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APPLICANT: Turn
                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 64:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 85.
Local Similarity nes 6; Conserv
                                                                         TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PETENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: GI 5190 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Megakaryocyte Stimulating Factors
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139 WMLSPFS 145
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                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                            LENGTH: 372 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                        NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 07/390,901 FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/457,196 FILING DATE: 29-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02140
                                                                                                                                                                                           TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
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Conservative
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Clark, Stephen C.
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                86.8%;
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 0; Mismatches
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Pred. No.
                Score 33; DB 6;
Pred. No. 5e+02;
                                                                                64:
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                               DB 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Version #1.25
                               Length 372;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
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   Indels
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TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-124-557-68
                                                                               RESULT 19
US-11-169-232-68
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US-10-124-557-68
  Sequence 68, Application US/11169232
Publication No. US20060025570A1
GENERAL INFORMATION:
APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 68, Application US/10124557 Publication No. US20020137894A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 68:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 422 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                              189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      139
                                                                                                                                                                                                    1 WMLSAFS 7
                                                                                                                                                                                                                                            6;
                                                                                                                                                            WMLSPFS 195
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                            Conservative
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Hewick, Rodney M.
Gesner, Thomas G.
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                                                                                                                                                                                                                                                          86.8%;
                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                            Score 33; DB 4;
Pred. No. 5.6e+02;
                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                            68:
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                                                                                                                                                                                                                                                                             Length 422;
                                                                                                                                                                                                                                          0
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US-10-124-557-66
; Sequence 66, Application US/10124557
; Publication No. US20020137894A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/446,114
FILING DATE: 29-UN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity hes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARBE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark, Stephen C.
Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetic
STREET: 87 Cambrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                          TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                          APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                            189 WMLSPFS 195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NABI: Cserr, Luann
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: GI 5190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/11/169,232 FILING DATE: 28-Jun-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 422 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
Genetics Institute,
CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                            86.8%;
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Pred. No. 5.6e+02;
0; Mismatches 1
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                    Inc.
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Length 422; Indels

0

CITY: Cambridge STATE: Massachusetts

Cambridge

FILING DATE: 28-Jun-2005

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US-11-169-232-66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 66, Application US/11169232
Publication No. US20060025570A1
GENERAL INFORMATION:
APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 66: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 66:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        190 WMLSPFS 196
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FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                 CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: «Unknown»
                                                                                                                                            COUNTRY: U.S.A.
ZIP: 02140
                                                                                                                                                                                                                           STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
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APPLICATION NUMBER: US/11/169,232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 423 amino acids
                                                                                                                                                                                                                                                 ADDRESSEE: Genetics Institute, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 5.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 86.8
Best Local Similarity 85.7
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 66:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/546,114
PILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 66:
                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          190 WMLSPFS 196
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NAME: Ceerr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                             STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Genetics Institute, STREET: 87 CambridgePark Drive CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                    APPLICATION NUMBER: US 07/643,502 FILING DATE: 18-JAN-1991 APPLICATION NUMBER: US 07/546,114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 423 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: <Unknown>
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/10124557
No. US20020137894A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gesner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hewick, Rodney M. Gesner, Thomas G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86.8%;
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Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 423;
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LENGTH: 463 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 54:
US-10-124-557-54
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                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Oun-2005
CLASSIFICATION NUMBER: US/11/169,232
FILING DATE: 28-Oun-2005
CLASSIFICATION OBTA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US/07/643,502
APPLICATION NUMBER: US 07/643,502
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-OAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/5457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
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Local Similarity 85.7%;
hes 6; Conservative
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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  TELECOMMUNICATION INFORMATION: TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           230 WMLSPFS 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WMLSAFS 7
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REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                           REGISTRATION NUMBER: 31,822 REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Genetics Institute, Inc. STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
TELEPHONE:
                                                                                       NAME: Cserr, Luann
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Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                               Sequence 27, Application US/09802207
Publication No. US20020086824A1
                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SU, JUI-LAN
TITLE OF INVENTION: SUPERFICIAL ZONE PROTEIN AND METHODS
TITLE OF INVENTION: MAKING AND USING SAME
FILE REFERENCE: 07083.0008U5
CURRENT APPLICATION NUMBER: US/10/038,694
CURRENT FILING DATE: 2001-12-31
PRIOR APPLICATION NUMBER: 60/258,920
PRIOR FILING DATE: 2000-12-29
APPLICANT: Carpten, John
APPLICANT: Trent, Jeffrey
APPLICANT: Trent, Jose
APPLICANT: Marcelino, Jose
TITLE OF INVENTION: Novel Methods and Reagents for the Treatment of Osteoarthritis
FILE REFERENCE: Case-06212
CURRENT APPLICATION NUMBER: US/09/802,207
CURRENT FILING DATE: 2001-08-29
PRIOR APPLICATION NUMBER: 09/619,175
PRIOR APPLICATION NUMBER: 60/145,328
PRIOR APPLICATION NUMBER: 60/145,328
                                                                                                                                                                           APPLICANT: Warman, Matthew APPLICANT: Carpten, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 54:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence; note OTHER INFORMATION: synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   305 WMLSPFS 311
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Kuettner, Klaus E.
Schmid, Thomas M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 463 amino acids TYPE: amino acid
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Schumacher, Barbara L.
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Pred. No. 6
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Pred. No. 6.9e+02;
0; Mismatches 1
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; TYPE: PRT
; ORCANISM: Homo sapiens
; FEATURE:
; NAME/KEY: NON CONS
; LOCATION: (267)...(268)
; NAME/KEY: NON CONS
; LOCATION: (321)...(322)
US-09-802-207-27
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US-10-425-115-337015
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US-10-124-557-44
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                                                                                                                                                                  Sequence 44, Application US/10124557 Publication No. US20020137894A1 GENERAL INFORMATION:
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SEQ ID NO 337015
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Publication No. US20040214272A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 30
SOFTWARE: PatentIn version
SEQ ID NO 27
                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (53222) B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: unsure
LOCATION: (1)..(1046)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
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Local Similarity 85.7%;
les 6; Conservative
                                                                                                                                                                                                                                                                                                                                                      Local Similarity 71.
                              TITLE OF INVENTION: Megakaryocyte
NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                    APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                     890 WLISAFS 896
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                                                                                                                                                                                                                                                                                                                      1 WMLSAFS 7
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ADDRESSEE: Genetics Institute, Inc
STREET: 87 CambridgePark Drive
                                                                                 Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                Clark, Stephen C.
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Pred. No. 9.7e+02;
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Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                   Stimulating
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RESULT 28
US-11-169-232-44
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Matches 6; Conserv
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: CBETY, LUANN
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: G1 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEPHONE: (617)876-1170
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 44:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE; Floppy disk
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1037 WMLSPFS 1043
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 WMLSAFS 7
                                                                                                                                                                                     STREET: 87 CambridgePark Drive
CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-CAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
                                                                                                                                                    COUNTRY: U.S.A. ZIP: 02140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                   ADDRESSEE: Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 1270 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                     Inc.
                                           Version #1.25
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US-10-124-557-42
Sequence 42, Application US/10124557
Publication No. US20020137894A1
GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-App-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAW-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                           COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-Jun-2005
CLASSIFICATION: <Unknown
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1037 WMLSPFS 1043
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                                                                                                                                                                                                                                                                                                                                                    STREET: 87 CambridgePark Drive
CITY: Cambridge
STATE: Massachusette
COUNTRY: U.S.A.
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
                                                                                                                                                                                                                                                                                                                                   ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 1270 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hewick, Rodney M. Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jacobs, Kenneth
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85.7%;
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Pred. No. 1.5e+03;
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                                                                                                                                                                                                                               Version #1.25
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RESULT 30
US-11-169-232-42
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Publication No.
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INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 1311 amino acids
                                                                                                                                                                                CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
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Local Similarity 85.7%;
hes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: CSerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 07/390,901 FILING DATE: 08-AUG-1989 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/11/169,232
FILING DATE: 28-Jun-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                    TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1078 WMLSPFS 1084
                                                                                                                       ATTORNEY/AGENT INFORMATION:
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Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Megakaryocyte Stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 WMLSAFS 7
                                                      NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 87 CambridgePark Drive
                                                                                                                                             APPLICATION NUMBER: US 07/390,901 FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (617)876-1170
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No. US20060025570A1
                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
E: (617)876-1170
(617)876-5851
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 33; DB 4; : Pred. No. 1.5e+03; 0; Mismatches 1
                                                            5190
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INFORMATION FOR SEQ ID NO: 42:

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TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 142:
US-10-124-557-142
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US-10-124-557-142
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Publication No. US20020
GENERAL INFORMATION:
APPLICANT: Turn
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Best Local Similarity
Query Match
Best Local Similarity
                                                                                                                                                                         REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-511
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 142:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
APTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 1311 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Genetics Institute, Inc. STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 31,822
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Cambridge
                                                                                                                                                        LENGTH: 1313 amino acids
                                                                                                                                                                                                                                                                                                                 NAME: Cserr,
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o. US20020137894A1
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86.8%;
                                                                                                                                                                                                                                                                                                                     Luann
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Pred. No. 1.5e+03;
Score 33; DB 4; Length 1313; Pred. No. 1.5e+03;
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TOPOLOGY: linear

HOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 142:
US-11-169-232-142
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US-11-169-232-142
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                                                                    Matches
                                                                                                  Query Match
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                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 142: SEQUENCE CHARACTERISTICS:
                                                                  Local Similarity les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
CLASSIFICATION: -Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
1080 WMLSPFS 1086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1080 WMLSPFS
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                                   1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
                                                                                                                                                                                                                                                                                                                               NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                                                                          LENGTH: 1313 amino acids
                                                                                                                                                                                                                                                                            TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 87 CambridgePark Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                  Conservative
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Clark, Stephen C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hewick, Rodney M.
Gesner, Thomas G.
                                                                                   86.8%;
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                                                                                   Score 33; DB 6;
Pred. No. 1.5e+03
                                                                      Mismatches
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                                                                                     .5e+03;
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                                                                                                     Length 1313;
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RESULT 33 US-10-124-557-50

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RESULT 34
US-11-169-232-50
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                                                                                                                                                                          Sequence 50, Application US/11169232 Publication No. US20060025570A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 50: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION ET : 16-Apr-2002
CLASSIFICATION ET : 16-Apr-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 86.8%;
Local Similarity 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                           APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                 1081 WMLSPFS 1087
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Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                          TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
        CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Cserr,
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                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                 Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
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Pred. No. 1.5e+03;
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US-10-124-557-46
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Publication No. US20020137894A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (617)876-5851 INFORMATION FOR SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION: «Unknown»

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/10/124,557

FILING DATE: 16-App-2002

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

ATTORNEY/AGENT INFORMATION:

NAME: CBETT, LUANIN

REFERENCE/DOCKET NUMBER: GI 5190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity es 6; Conserv
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
                                                                                                                                                                                                                 TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1081 WMLSPFS 1087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION
                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetic
                                                                                                                                                                                                                                                                                                                        APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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                                                                   ZIP:
                                                                                     CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                  ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
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CITY: Cambridge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH: 1314 amino acids
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                                                                                                                                                                                                                                                           Hewick, Rodney M. Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                      Jacobs, Kenneth
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Pred. No. 1.5e+03;
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Length 1314; Indels

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US-10-124-557-60
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Publication No. US20020137894A1
GENERAL INFORMATION:
APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 1320 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 46:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                 ZIP: 02140
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-App-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1087 WMLSPFS 1093
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
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FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
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REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: < Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 16-Apr-2002
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                                                                                                                                                                                                                                                                                                                                    CITY: Cambridge
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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Clark, Stephen C.
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Pred. No. 1.5e+03;
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US-11-169-232-46
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Publication No. US20050106579A1
GENERAL INFORMATION:
                                                                                                                                                                                      Sequence 46, Application US/11169232 Publication No. US20060025570A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1
SEQ ID NO 58
LENGTH: 1320
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 86.8%;
Best Local Similarity 85.7%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: OriGene Technologies, Inc

TITLE OF INVENTION: Regulated Angiogenesis Genes and Polypeptides

FILE REPERENCE: 1U 103 R1

CURRENT APPLICATION NUMBER: US/10/717,665

CURRENT FILING DATE: 2003-11-21

PRIOR APPLICATION NUMBER: US/10/164,595

PRIOR APPLICATION NUMBER: US/10/164,595

PRIOR FILING DATE: 2002-06-10

NUMBER OF SEQ ID NOS: 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617)876-1170
TELEPAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1087 WMLSPFS 1093
                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetic
                                                        TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                              1087 WMLSPFS 1093
                                                                                                                                                  APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Cserr, Luann
NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
REGISTRATION NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 1320 amino acids
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                         Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                           86.8%;
                  Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Pred. No. 1.5e+03;
0; Mismatches 1
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Pred. No. 1.5e+03;
0; Mismatches 1;
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                    Inc.
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RESULT 39
US-11-169-232-60
US-11-169-232-60; Sequence 60, Application US/11169232; Publication No. US20060025570A1; GENERAL INFORMATION:
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear;

MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
US-11-169-232-46
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 46: SEQUENCE CHARACTERISTICS:
ZIP: 0Z14v
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Unn-2005
CLASSIFICATION: CUnknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UNN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                       Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
                                                                                                                                                                       CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                               ADDRESSEE: Genetics Institute, Inc
STREET: 87 CambridgePark Drive
                                                                                                                                                      COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH: 1320 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86.8%;
85.7%;
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                                 Version #1.25
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US-10-124-557-48
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Publication No. US20020137894A1
GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 85.
                                                                                                                                          COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ceerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
                               CLASSIFICATION: <Ünknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 60:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Megakaryocyte Stimulating Factors NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1087 WMLSPFS 1093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                  STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/643,502 FILING DATE: 18-JAN-1991 APPLICATION NUMBER: US 07/546,114 FILING DATE: 29-UN-1990 APPLICATION NUMBER: US 07/457,196 FILING DATE: 29-DEC-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/11/169,232 FILING DATE: 28-Jun-2005
                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Genetics Institute, STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                         CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 1320 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86.8%;
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Pred. No. 1.5e+03;
0; Mismatches 1
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Gaps

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RESULT 41
US-11-169-232-48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 48, Application US/11169232 Publication No. US20060025570A1 GENERAL INFORMATION:
APPLICANT: Turner, Katherine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 48: SEQUENCE CHARACTERISTICS:
                                                                             APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 86.8%;
Local Similarity 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: CSETY, LUANN
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
TELEPHONE: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 02140
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 48:
                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI 5190
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1121 WMLSPFS 1127
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FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 87 CambridgePark Drive
                                                                   NAME: Cserr,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1354 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Clark, Stephen C.
                                                                 Luann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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Pred. No. 1.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Query Match
Best Local Similarity
Watches 6; Conserva
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US-10-124-557-40
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US-10-124-557-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 40, Application US/10124557 Publication No. US20020137894A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 48: SEQUENCE CHARACTERISTICS:
                                                                                                               TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
              TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 40:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1121 WMLSPFS 1127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 48:
                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS: ADDRESSE: Genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 WMLSAFS 7
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                            NAME: Cserr, Luann
NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linea:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                           TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 1354 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX:
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                                                                                                ENGTH: 1361 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jacobs, Kenneth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genetics Institute,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 33; DB 6;
Pred. No. 1.6e+03;
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RESULT 43
US-11-169-232-40
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Best Local S
Matches 6
                                                                                   Query Match
Best Local Similarity
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Publication No. US2006
GENERAL INFORMATION:
                                                                      Matches
                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Un-2005
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Turner, Katherine
Clark, Stephen C.
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1128 WMLSPFS 1134
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                                   1 WMLSAFS 7
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                                                                                                                                                                                                                                                                                                                                    NAME: Cserr, Luann
NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                            TYPE: amino acid
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                                                                                                                                                                                                                            LENGTH: 1361 amino acids
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                                                                                                                                                                                                                                                                                  TELEFAX: (617)876-5851
                                                                    Conservative
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85.7%;
                                                                                   86.8%;
85.7%;
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                                                                                   Score 33; DB 6;
Pred. No. 1.6e+03;
                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                        5190
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                                                                    1; Indels
                                                                                                    Length 1361;
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RESULT 44
US-10-124-557-52
Sequence 52, Application US/10124557
Publication No. US20020137894A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-124-557-52
                                                                                                                                                                  RESULT 45
US-11-169-232-52
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                                                                                                          Sequence 52, Application No. US2006 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 29-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compartible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
             APPLICANT: Turner, Katherine
Clark, Stephen C.
Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
                                                                                                                                                                                                                                               1130 WMLSPFS 1136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: GI 5190 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Turner, Katherine
Clark, Stephen C.
TITLE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Megakaryocyte Stimulating Factors
                                                                                                                                                                                                                                                                                   1 WMLSAFS
                                                                                                                                                                                                                                                                                                                       6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/10/124,557 FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                Application US/11169232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 1363 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
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                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                            US20060025570A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (617)876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genetics Institute,
                                                                                                                                                                                                                                                                                                                                          86.8%;
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Megakaryocyte Stimulating Factors
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Pred. No. 1.6e+03;
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                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                        Sequence 30, Application US/09802207 Publication No. US20020086824A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                    CURRENT APPLICATION NUMBER: US/09/802,207
CURRENT FILING DATE: 2001-08-29
PRIOR APPLICATION NUMBER: 09/619,175
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 60/145,328
PRIOR FILING DATE: 1999-07-23
                                                                                                                                                            APPLICANT: Warman, Matthew
APPLICANT: Carpten, John
APPLICANT: Trent, Jeffrey
APPLICANT: Trent, Jose
APPLICANT: Marcelino, Jose
TITLE OF INVENTION: Novel Methods and Reagents for the Treatment of Osteoarthritis
FILE REFERENCE: Case-06212
NUMBER OF SEQ ID NOS: 30 SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/11/169,232
FILING DATE: 28-Jun-2005
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                  1130 WMLSPFS 1136
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 WMLSAFS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/10/124,557
FILING DATE: 16-Apr-2002
APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-ANN-191
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-UN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Cserr, Luann
REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Cambridge
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1363 amino acids
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Pred. No. 1.6e+03;
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Query Match
Best Local Similarity
Matches 6; Conserve
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Best Local Similarity by...
Grandes 6; Conservative
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Publication No. US20040072741A1
GENERAL INFORMATION:
APPLICANT: Jay, Gregory D.
TITLE OF INVENTION: Tribonectin Po
                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10124557 Publication No. US20020137894A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/897,188
CURRENT FILING DATE: 2001-07-02
PRIOR APPLICATION NUMBER: 09/298,970
PRIOR FILING DATE: 1999-04-23
PRIOR APPLICATION NUMBER: 09/556,246
PRIOR APPLICATION NUMBER: 09/556,246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 2000-04-24
NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: 21486-026 CIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 1404
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1404
 ZIP: 02140

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/124,557

FILING DATE: 16-Apr-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1171 WMLSPFS 1177
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                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Megakaryocyte Stimulating NUMBER OF SEQUENCES: 143
                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Turner, Katherine
Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 WMLSAFS 7
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                                                                                                                                                                              STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                   ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                     CITY: Cambridge
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                               Hewick, Rodney M. Gesner, Thomas G.
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Pred. No. 1.6e+03;
0; Mismatches
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Pred. No.
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TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-124-557-2
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US-10-124-557-62
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Best Local Similarity 85.7%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 2:
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/124,557

FILING DATE: 16-Apr-2002

CLASSIFICATION: CURROWN>
PRIOR APPLICATION: CURROWN>
PRIOR APPLICATION NUMBER: US 07/643,502

FILING DATE: 18-JAN-1991

APPLICATION NUMBER: US 07/546,114

FILING DATE: 18-JUN-1990

APPLICATION NUMBER: US 07/457,196

FILING DATE: 29-DEC-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

APPLICATION NUMBER: US 07/390,901

FILING DATE: 08-AUG-1989

APPLICATION NUMBER: US 07/390,901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/643,502
FILING DATE: 18-JAN-1991
APPLICATION NUMBER: US 07/546,114
FILING DATE: 29-JUN-1990
APPLICATION NUMBER: US 07/457,196
FILING DATE: 29-DEC-1989
APPLICATION NUMBER: US 07/390,901
FILING DATE: 08-AUG-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                              ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: ISM PC TOOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jacobs, Kenneth
Hewick, Rodney M.
Gesner, Thomas G.
TITLE OF INVENTION: Megakaryocyte Stimulating Factors
NUMBER OF SEQUENCES: 143
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TELEPHONE: (617)876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Turner, Katherine Clark, Stephen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1404 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10124557
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; NUMBER OF SEQ ID NOS: 16
; SOFTMARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 1404
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-858-595-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: protein ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-124-557-62
Search completed: August 29, 2006, 10:18:50 Job time : 87.6931 secs
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Best Local Similarity
Matches 6; Conserv
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PRIOR APPLICATION NUMBER: US/09/298,970
PRIOR FILING DATE: 1999-04-23
                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/858,595
CURRENT FILING DATE: 2004-06-01
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Jay, Gregory D.
TITLE OF INVENTION: TRIBONECTINS
FILE REFERENCE: 21486-026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (617)876-5851
INFORMATION FOR SEQ ID NO: 62:
SEQUENCE CHARACTERISTICS:
LENGTH: 1404 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
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85.7%;
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Pred. No. 1.6e+03;
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Pred. No. 1.6e+03;
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Result
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Maximum
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Perfect score:
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Match
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Gapop 10.0 ,
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38
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US-10-449-902-38264
US-11-174-307B-5396
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US-11-353-390A-14
US-11-354-310A-14
US-10-449-902-52867
US-10-449-902-53284
US-10-449-902-53284
US-10-953-349-38417
US-11-056-355B-82525
US-10-511-455-26
US-11-056-355B-82525
US-10-953-349-3329
US-11-056-355B-82529
US-10-953-349-38416
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US-10-953-349-38416
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US-10-953-349-38416
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US-10-953-349-38416
US-11-056-355B-82529
US-10-953-349-38416
US-11-056-355B-179382
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US-11-056-355B-117746
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Biocceleration
 Sequence 28641,
Sequence 396, A;
Sequence 14, App
Sequence 14, App
Sequence 14, App
Sequence 14, App
Sequence 52867,
Sequence 53284,
Sequence 338417,
Sequence 79383,
Sequence 26, App
Sequence 26, App
Sequence 3329,
Sequence 31231,
Sequence 11231,
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  e 14, Appl

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e 52857, A

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US-110-953-349-33297
US-110-953-349-33297
US-110-953-349-33297
US-110-953-349-31756
US-110-953-349-31829
     sequence
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75-11-024-544A-16
75-11-024-737-120
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IS-10-449-902-37851
IS-11-049-902-37851
IS-11-056-355B-9999
IS-11-330-403-1369
IS-11-330-403-1369
IS-11-330-403-355B-9105
IS-11-330-403-355B-9105
IS-11-056-355B-90208
IS-11-10-65-355B-90208
IS-11-056-355B-90208
IS-11-056-355B-90208
IS-11-30-409-902-40950
IS-11-30-409-902-40950
IS-11-30-409-902-40950
IS-11-056-355B-90306
IS-11-056-355B-90306
IS-11-056-355B-90308
                                                                                                                  S-10-528-872-3

S-10-449-902-6608

S-11-330-403-4196

S-11-056-3558-80481

S-11-056-3558-86416

S-11-221-332-12

S-11-221-332-12
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Sequence
921199, p. 231006, p. 
                                                      GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: National Institute of Physical and Chemical Research.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
FRIOR APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
FRIOR APPLICATION NUMBER: JP 2002-203269
FRIOR APPLICATION NUMBER: JP 2002-383870
FRIOR APPLICATION NUMBER: JP 2002-383870
FRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 28641
FILE DATE: 2012-05-10
GRGANISM: Oryza sativa
US-10-449-902-28641
                                                                                                                                                                                                                                                                                                           US-10-449-902-28641

Sequence 28641, Application US/10449902

Publication No. US20060123505A1

GENERAL INFORMATION:
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                                   Query Match
Best Local S
Matches
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                                   . Similarity 5; Conserv
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US-11-374-317-174-3
US-11-274-317-174-3
US-11-241-596-96
US-11-241-596-96
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-248-718-105
US-11-249-902-41029
US-11-449-902-55047
US-11-245-628-43
US-11-174-3078-3672
US-11-174-3078-368-90300
US-11-174-3078-3588-90300
US-11-056-3558-90300
                                   Score 31; DB Pred. No. 61; 1; Mismatches
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273, Appl
17433, A
966, A37, A
966, A37, A
93962, Appl
105, App
1
                                    Gaps
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PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 38264
LENGTH: 302
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US-10-449-902-38264
RESULT 4
US-11-399-675-14
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US-11-174-307B-5396
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Best Local
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LENGTH: 302
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Publication No. US20060123505A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/583,671
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,781
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,651
PRIOR FILING DATE: 2004-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: UCCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY
TITLE OF INVENTION: USEFUL FOR MODIFYING PLANT CHARACTERISTICS
FILE REFERENCE: 2750-1601PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/11/174,307B
CURRENT FILING DATE: 2005-06-30
                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
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Similarity 71.4%;
5; Conservative
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5; Conserva
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Pred. No. 62;
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Pred. No. 62;
1; Mismatches
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US-11-354-310A-14

Sequence 14, Application US/11354310A Publication No. US20060185037A1 GENERAL INFORMATION: APPLICANT: Sticklen, Masomeh B

APPLICANT:

Masomeh Shahina

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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                            SOFTWARE: PatentIn version 3.3 SEQ ID NO 14
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Publication No. US20060185036A1
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SEQ ID NO 14
LENGTH: 371
TYPE: PRT
ORGANISM: Phanerochaete chrysosporium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14, Application US/11399675 Publication No. US20060179513A1 GENERAL INFORMATION:
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Matches 6; Conserv
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CURRENT FILING DATE: 2006-04-06
PRIOR APPLICATION NUMBER: US 60/242,408
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 09/981,900
PRIOR PILING DATE: 2001-10-18
NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sticklen, Masomeh B
APPLICANT: Magbool, Shahina B
APPLICANT: Magbool, Shahina B
APPLICANT: Dale, Bruce E
TITLE OF INVENTION: TRANSGENIC PLANTS CONTAINING LIGNINASE AND CELLULASE WHICH
TITLE OF INVENTION: DEGRADE LIGNIN AND CELLULOSE TO FERMENTABLE SUGARS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Magbool, Shahina B
APPLICANT: Magbool, Shahina B
APPLICANT: Dale, Bruce E
TITLE OF INTERPRETATION
                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 09/981,900
PRIOR FILING DATE: 2001-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TRANSGENIC PLANTS CONTAINING LIGNINASE AND CELLULASE TITLE OF INVENTION: DEGRADE LIGNIN AND CELLULOSE TO FERMENTABLE SUGARS FILE REFERENCE: MSU 4.1-814
                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: MSU 4.1-806
CURRENT APPLICATION NUMBER: US/11/353,390A
CURRENT FILING DATE: 2006-02-14
CURRENT APPLICATION NUMBER: US 60/242,408
PRIOR APPLICATION NUMBER: US 60/242,408
                                                                                                                                                                               TYPE: PRT ORGANISM: Phanerochaete chrysosporium
                                                                                                                                                                                                                      ENGTH:
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197 WMLSAHS 20
197 WMLSAHS 203
                       1 WMLSAFS 7
                                                                               Conservative
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85.7%;
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                                                                             Score 31; DB Pred. No. 75; O; Mismatches
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Pred. No.
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT APPLICATION NUMBER: US/20/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-338870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR APPLICATION NUMBER: JP 2002-383870
PRIOR PRIOR DATE: 2002-12-11
NUMBER OF SEQ ID MOS: 56791
SOFTWARE: Patentin Ver. 2.1
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GENERAL INFORMATION:

APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF

FILE REFERENCE: MOA-A0205Y1-US

CURRENT APPLICATION NUMBER: US/10/449,902
                                                                                                                                                                                           Sequence 46155, Application US/10449902 Publication No. US20060123505A1 GENERAL INFORMATION:
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LENGTH: 504
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Best Local Similarity 85.7%;
Matches 6; Conservative
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-10-449-902-52867
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CURRENT APPLICATION NUMBER: US/11/354,310A
CURRENT FILING DATE: 2006-02-14
PRIOR APPLICATION NUMBER: US 60/242,408
PRIOR PILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 09/981,900
PRIOR APPLICATION NUMBER: US 09/981,900
PRIOR FILING DATE: 2001-10-18
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TYPE: PRT
ORGANISM: Phanerochaete chrysosporium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 71.
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71.4%;
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Pred. No. 1.6e+02;
1; Mismatches 1
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Pred. No. 75;
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; ORGANISM: Oryza sativa
US-10-449-902-53284
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US-10-449-902-46155
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SEQ ID NO 53284
LENGTH: 63
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PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46155
LENGTH: 1128
                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                           Sequence 44647, Application US/10449902
Publication No. US20060123505A1
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APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/SCURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: UF 2002-203269
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-12-11
ROMEREN OF SEQ ID NOS: 56791
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SOFTWARDER OF SEC ID NOS: 56791
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Matches 5; Conserv
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Best Local Similarity
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APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-0US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-05-30
PRIOR FILING DATE: 2002-12-11
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PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
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Pred. No. 34;
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; FEATURE:
; NAME/KEY: peptide
; LOCATION: (1)...(297)
; OTHER INFORMATION: Ceres Seq. ID no. 12651075
US-11-056-355B-79384
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US-10-953-349-38417
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US-11-056-355B-79384
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; ORGANISM: Oryza sativa
US-10-449-902-44647
                                                                                       , ORGANISM: Zea mays subsp. mays
US-10-953-349-38417
                                                                                                                   Sequence 38417, Application US/10953349

Publication No. US20060107345A1

GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT EPLING DATE: 2004-09-30
RUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 38417
LENGTH: 308
TYPE: PRT
Query Match
Best Local Similarity 71.7
Conservative
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 44647
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Best Local :
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SEQ ID NO 79384
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Best Local Similarity 57.1%;
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ORGANISM: Arabidopsis thaliana
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                                    76.3%;
                      Score 29; DB 6; Le
Pred. No. 1.6e+02;
1; Mismatches 1;
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Pred. No. 1.5e+02;
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Pred. No. 51;
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US-11-056-355B-79383
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APPLICANT: Alexandrov, Nickolai
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590FUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 82555
LENGTH: 334
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FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 79383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 82525, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
                                                                                                                                   TYPE: prt
ORCANISM: Arabidopsis thaliana
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(334)
OTHER INFORMATION: Ceres Seq. II
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Matches 5; Conserv
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                                                                             Query Match
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LOCATION: (1)..(308)
OTHER INFORMATION: Ceres Seq. ID no. 12651074
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ORGANISM: Arabidopsis thaliana
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1 WMLSAFS 7
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57.1%;
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                                                            Score 29; DB 7;
Pred. No. 1.7e+02;
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Pred. No. 1.6e+02;
1; Mismatches 1
                                          Mismatches
                                                                                                                                            no. 12666048
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US-10-511-455-26

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PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: GB0207904.4
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: GB0207900.2
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: GB0207901.0
PRIOR APPLICATION NUMBER: GB0207901.0
PRIOR APPLICATION NUMBER: GB0227734.1
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: GB0227734.1
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Publication No. US20060088835A1
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                                                             SEQ ID NO 4
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Publication No. US20060177904A1
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APPLICANT:
APPLICANT:
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                                                                            APPLICANT: Fechtel, Kim
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 00766,000103.6
CURRENT APPLICATION NUMBER: US/11/395,249
CURRENT FILING DATE: 2006-04-03
NUMBER OF SEQ ID NOS: 240
NUMBER OF SEQ ID NOS: 240
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APPLICANT: McCoy,
APPLICANT: LaVall:
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SOFTWARE: PatentIn versi
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CURRENT FILING DATE: 2004-10-05
PRIOR APPLICATION NUMBER: PCT/GB03/001543
PRIOR FILING DATE: 2003-04-07
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TITLE OF INVENTION: SCHIZOPHRENIA ASSOCIATED GENES
                                                                                                                                                                                                                                                                                     APPLICANT:
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ORGANISM: Homo sapiens
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                                        ENGTH: 335
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Local Similarity 83.3%;
es 5; Conservation
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                                                                                                                                                                                                                   Agostino, Michael J.
Steininger II, Robert J.
Spaulding, Viki
Wong, Gordon G.
Clark, Hilary
                                                                                                                                                                                                                                                                                                                                                                                                           McCoy, John M.
LaVallie, Edward R.
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                                                                                                                                                                                                                                                                                                                           Merberg, David
Treacy, Maurice
                                                                                                                                                                                                                                                                                                                                                                                    Collins-Racie, Lisa
                                                                                                                                                                                                                                                                                                                                                                   Evans, Cheryl
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Pred. No. 1.
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1.7e+02;
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Query Match
Best Local Similarity
Watches 5; Conserve
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US-10-953-349-33299
; Sequence 33299, Application US/10953349
; Publication No. US20060107345A1
; GENERAL INFORMATION:
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                                                     Query Match
Best Local Similarity
Matches 5; Conserve
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US-11-056-355B-82529
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                                                                                                                                                                        APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT PILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 33299
LENGTH: 345
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding TITLE OF INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2 CURRENT APPLICATION NUMBER: US/11/056,3558 CURRENT FILING DATE: 2005-02-14 PRIOR APPLICATION NUMBER: 60/544,190 PRIOR FILING DATE: 2004-02-13 NUMBER OF SEQ ID NO 8259 LENGTH: 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 82529, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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Best Local (
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LOCATION: (1)..(338)
OTHER INFORMATION: Ceres Seq. ID no. 12666052
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ORGANISM: Arabidopsis thaliana
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nes 4; Conserv
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310 WLLAAFT 316
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270 WLLIAFS
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83.3%;
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57.1%;
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    Mismatches

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                                                                                      Score 29; DB 6;
Pred. No. 1.8e+0
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Pred. No. 1.
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Pred. No. 1.7e+02;
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                                                                                       .8e+02;
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APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
                                                                                                                                 Sequence 30511, Application US/10449902 Publication No. US20060123505A1 GENERAL INFORMATION:
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Best Local Similarity 71.4%;
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CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
APPLICANT: National Institute of Agrobiological Sciences.
APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Paylical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding TITLE OF INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2 CURRENT APPLICATION NUMBER: US/11/056,355B CURRENT FILING DATE: 2005-02-14
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TYPE: prt
ORGANISM: Arabidopsis thaliana
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: National Institute of Agrobiological Sciences.

APPLICANT: Bio-oriented Technology Research Advancement Institution.

APPLICANT: The Institute of Physical and Chemical Research.

APPLICANT: Foundation for Advancement of International Science.

ITILE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF FILE REFERENCE: MOA-A0205Y1-US

CURRENT FILING DATE: 2003-05-29

PRIOR APPLICATION NUMBER: US/10/449,902

CURRENT FILING DATE: 2002-05-30

PRIOR APPLICATION NUMBER: UP 2002-203269

PRIOR APPLICATION NUMBER: UP 2002-383870

PRIOR APPLICATION NUMBER: JP 2002-383870

PRIOR FILING DATE: 2002-12-11

NUMBER OF SEG ID NOS: 56791

SOFTWARE: PatentIn Ver. 2.1

SOFTWARE: PatentIn Ver. 2.1
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US-10-449-902-49957
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US-10-449-902-30511
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Sequence 32815, Application US/10953349
Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, NICKOLAI et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTMARE: PATENTIN Version 3.3
SEQ ID NO 32815
LENGTH: 355
TYPE: PRT
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SEQ ID NO 30511
LENGTH: 354
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Publication No. US20060123505A1
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CURRENT FILLING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: JP 2002-203269
PRIOR FILING DATE: 2002-05-30
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les 5; Conserv
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71.4%;
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    Mismatches

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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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; FEATURE:
NAME/KEY: peptide
; LOCATION: (1).. (355)
; OTHER INFORMATION: Ceres Seq. ID no. 13492044
US-11-056-355B-17746
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US-11-056-355B-17746
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                                                                                                                                                                  TITLE OF INVENTION: Sequence Determined DNA Fragments of INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 17746
LENGTH: 355
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APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
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Query Match
Best Local
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NAME/KEY: peptide
LOCATION: (1)..(355)
OTHER INFORMATION: Ceres Seq. ID no.
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ORGANISM: Zea mays
                                                                                                                                  ORGANISM: Zea mays subsp. mays
                                                                                                                                                     TYPE: prt
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Similarity
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Pred. No. 1.8e
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Pred. No. 1.8e+02;
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DB 7;
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APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA
TITLE OF INVENTION: ENCOMDED THERBY
FILE REPERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 33298
LENGTH: 357
                                                                                  US-11-330-403-6594
US-11-330-403-6594
; Sequence 6594, Application US/11330403
; Publication No. US20060159563A1
; GENERAL INFORMATION:
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; ORGANISM: Zea mays subsp. mays
US-10-953-349-33298
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US-10-953-349-38415
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Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DN
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-15799US2
CURRENT APPLICATION NUMBER: US/10/953,349
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Publication No. US20060107345A1
GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3
SEQ ID NO 38415
LENGTH: 357
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Best Local
APPLICANT: Abad, Mark S.
TITLE OF INVENTION: Genes and Uses for Plant Improvement FILE REFERENCE: 38 8-21 (53629) B
CURRENT APPLICATION NUMBER: US/11/330,403
CURRENT FILING DATE: 2006-01-12
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Pred. No. 1.8e+02;
1; Mismatches 1
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Pred. No. 1.8e+02;
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Best Local Similarity
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US-10-953-349-33297
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US-10-449-902-38671
                                                                                    ; ORGANISM: Zea mays subsp. mays
US-10-953-349-33297
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Publication No. US20060107345A1
GENERAL INFORMATION:
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SEQ ID NO 6594
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                                           Query Match
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        Matches
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-15799US2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40052
SOFTWARE: PRESENTIN Version 3.3
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT cDNA AND USES THEREOF
FILE REFERENCE: MOA-A0205Y1-US
CURRENT APPLICATION NUMBER: US/10/449,902
CURRENT FILING DATE: 2003-05-29
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PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
SOFTWARE: Patentin Ver. 2.1
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TYPE: PRT
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                                                                                                                                           LENGTH: 417
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Local Similarity 71.4%;
nes 5; Conservativo
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83.3%;
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Pred. No. 2e+02;
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Pred. No. 1.9e+02;
0; Mismatches 1
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                                           Length 417;
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US-11-056-355B-17744, Application US/11056355B; Sequence 17744, Application US/11056355B; Publication No. US20060150283A1; GENERAL INFORMATION: Prover, Vyacheslav
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                                                                                                RESULT 33
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Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 32813
LENGTH: 418
TYPE: PRT
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CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 11229
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ORGANISM: Zea mays subsp. m
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(418)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Brover, Vyacheslav APPLICANT: Alexandrov, Nickolai TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding TITLE OF INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2
                                                                                                                                                                                                                                                                                                                              LOCATION: (1)..(418)
OTHER INFORMATION: C
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343 WLLIAFS 349
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71.4%;
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Pred. No.
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Pred. No. 2.1e+02
1; Mismatches
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RESULT 35
US-11-056-355B-82528
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; NAME/KEY: peptide
; LOCATION: (1)...(437)
; OTHER INFORMATION: Ceres Seq. ID no. 12666047
US-11-056-355B-82524
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Best Local Similarity
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TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT FILING DATE: 2005-02-14
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 82524
LENGTH: 437
                                                                                                                       Sequence 82528, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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Best Local
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                 APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590FUS2
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  CURRENT APPLICATION
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LOCATION: (1)..(418)
OTHER INFORMATION: C
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ORGANISM: Arabidopsis thaliana
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ORGANISM: Zea mays subsp. mays
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4; Conservative
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  NUMBER:
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57.1%;
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71.4%;
US/11/056,355B
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Pred. No. 2.2e+02
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2.1e+02
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US-11-056-355B-66755

Sequence 66755, Application US/11056355B

Publication No. US20060150283A1

GENERAL INFORMATION:

APPLICANT: Brover, Vyacheslav

APPLICANT: Alexandrov, Nickolai

TITLE OF INVENTION: Sequence Determined DNA Fragments and

TITLE OF INVENTION: Polypeptides Encoded Thereby

FILE REFERENCE: 2750-1590PUS2

CURRENT APPLICATION NUMBER: US/11/056,355B

CURRENT APPLICATION NUMBER: 60/544,190

PRIOR APPLICATION NUMBER: 60/544,190

PRIOR APPLICATION NUMBER: US/11/056,355B

CURRENT FILING DATE: 2004-02-13

NUMBER OF SEQ ID NOS: 119966

SEQ ID NO 66755

LENGTH: 495

TYPE: nort
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CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 31757
LENGTH: 495
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Publication No. US20060107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
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PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
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ORGANISM: Arabidopsis thaliana
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NAME/KEY: peptide
LOCATION: (1)..(438)
OTHER INFORMATION: Ceres Seq. ID no. 12666051
                                              TYPE: prt
ORGANISM: Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
NAME/KEY: peptide
LOCATION: (1)..(495)
                                        FEATURE:
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57.1%;
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71.4%;
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Pred. No. 2.
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Pred. No. 2.2e+02;
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2.5e+02;
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US-11-056-355B-66754
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APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOCTWARE: Patentin version 3.3
SEQ ID NO 31756
LENGTH: 530
                                                                                                                                                                                                    TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding TITLE OF INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2 CURRENT APPLICATION NUMBER: US/11/056,355B CURRENT FILING DATE: 2005-02-14 PRIOR APPLICATION NUMBER: 60/544,190 PRIOR FILING DATE: 2004-02-13 NUMBER OF SEQ ID NOS: 119966 SEQ ID NO 66754 LENGTH: 530 TYPE: prt ORGANISM: Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 66754, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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NAME/KEY: peptide
LOCATION: (1)..(530)
OTHER INFORMATION: Ce
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 WALAAFS 101
                                WMLSAFS 7
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Pred. No. 2.5e
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Pred. No. 2.7e+02;
1; Mismatches 1
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                                                                              Score 29; DB 7;
Pred. No. 2.7e+02
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2.5e+02;
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                                                                                             Length 530;
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US-11-056-355B-82527
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US-11-056-355B-82527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 82527, Application US/11056355B
Publication No. US20060150283A1
GENERAL INFORMATION:
APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 82523
LENGTH: 564
                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 82527
                                                                      Matches
                                                                                        Best
                                                                                                          Query Match
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                                                                                                                                                                      FEATURE:

NAME/KEY: peptide
LOCATION: (1)..(56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
NUMBER OF SEQ ID NOS: 119966
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding TITLE OF INVENTION: Polypeptides Encoded Thereby FILE REFERENCE: 2750-1590PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: peptide
LOCATION: (1)..(564)
OTHER INFORMATION: Ceres Seq. ID no. 12666046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: prt
ORGANISM: Arabidopsis thaliana
FEATURE:
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ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                               LENGTH: 565
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                                                                                        Local
537
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                                   1 WMLSAFS 7
                                                                  Similarity
4; Conserve
WLLAAFT 543
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                                                                      Conservative
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57.1%;
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57.1%;
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                                                                                        Score 29; DB 7;
Pred. No. 2.9e+02;
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Pred. No. 2.9e+02;
                                                                        Mismatches
                                                                                                          DB 7;
                                                                                                        Length 565;
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RESULT 42 US-10-449-902-47030 ; Sequence 47030, Application US/10449902 ; Publication No. US20060123505A1

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US-11-056-3558-66753
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US-10-953-349-31755
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                                                                                                                                                                                             Sequence 66753, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
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SEQ ID NO 31755
LENCTH: 610
TYPE: PRT
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Publication No. US20060107345A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2003-05-29
PRIOR APPLICATION NUMBER: UP 2002-203269
PRIOR FILING DATE: 2002-05-30
PRIOR APPLICATION NUMBER: UP 2002-383870
PRIOR FILING DATE: 2002-12-11
NUMBER OF SEQ ID NOS: 56791
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                                                                                   APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REPERBNCE: 2750-1579PUS2
CURRENT REPERBNCE: 2750-1579PUS2
CURRENT FILINGIOTION UMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
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-10-449-902-47030
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APPLICANT: Bio-oriented Technology Research Advancement Institution.
APPLICANT: The Institute of Physical and Chemical Research.
APPLICANT: Foundation for Advancement of International Science.
TITLE OF INVENTION: FULL-LENGTH PLANT CDNA AND USES THEREOF
FILE REFERENCE: MOA-A020571-US
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2005-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
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TYPE: PRT
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5; Conserv
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71.4%;
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; Pred. No. 3.1e.
1; Mismatches
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Pred. No. 3e+02;
1; Mismatches
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Query Match
Best Local Similarity
Matches 5; Conserve
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US-10-953-349-33830
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US-10-953-349-33831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT FILING DATE: W/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 33831
LENGTH: 638
TYPE: PRT
                                                                                                                                                                                                                         Sequence 33830, Application US/10953349
Publication No. US20066107345A1
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai et al.
AITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579US2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
CURRENT FILING DATE: 2004-09-30
                                                                                                                                                         NUMBER OF SEQ ID NOS: 40252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 33830
LENGTH: 665
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ORGANISM: Triticum aestivum
FEATURE:
NAME/KEY: peptide
LOCATION: (1)..(610)
OTHER INFORMATION: Ceres Seq. I
                                    Best Local Similarity
Matches 5; Conserv
                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                       ORGANISM: Zea mays subsp. mays
                                                                                                                                             TYPE:
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   1 WMLSAF 6
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83.3%;
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83.3%;
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71.4%;
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                                                      Score 29; DB 6;
Pred. No. 3.4e+02;
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Pred. No. 3.2e+02;
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Pred. No. 3.1e+02;
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                                      Mismatches
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                                                                        Length 665;
                                      Indels
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US-11-056-355B-1495
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APPLICANT: ALEXANDROV, Nickolai et al.
APPLICANT: ALEXANDROV, Nickolai et al.
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: ENCONDED THERBY
FILE REFERENCE: 2750-1579PUS2
CURRENT APPLICATION NUMBER: US/10/953,349
CURRENT FILING DATE: 2004-09-30
NUMBER OF SEQ ID NOS: 40252
SOFTWARE: Patentin version 3.3
SEQ ID NO 33829
LENGTH: 667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
Sequence 1495, Application US/11056355B
publication No. US20060150283A1
GENERAL INFORMATION:
APPLICANT: Brover, Vyacheslav
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments
TITLE OF INVENTION: Polypeptides Encoded Thereby
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SEQ ID NO 1496
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/544,190 PRIOR FILING DATE: 2004-02-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/11/056,355B
                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: peptide
LOCATION: (1)..(105)
OTHER INFORMATION: Ceres Seq.
                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: prt
ORGANISM: Zea mays subsp. mays
FEATURE:
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                                                                                                                                                                                                                                                                                           Local Similarity
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83.3%;
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Pred. No. 3.4e+02;
Pred. No. 3.4e+12;
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; FEATURE:
, NAME/KEY: peptide
; LOCATION: (1) . (154)
; OTHER INFORMATION: Ceres Seq.
US-11-056-355B-1494
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Search completed: August 29, Job time : 5.54455 secs
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                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 119966
SEQ ID NO 1494
SEQ ID: 154
TYPE: Drt
ORGANISM: Zea mays subsp. ma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1494, Application US/11056355B Publication No. US20060150283A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best
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LENGTH: 124
                                                                                                                                      Matches
                                                                                                                                                        Best
                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Brover, Vyacheslav
APPLICANT: Brover, Nickolai
APPLICANT: Alexandrov, Nickolai
TITLE OF INVENTION: Sequence Determined DNA Fragments and Corresponding
TITLE OF INVENTION: Polypeptides Encoded Thereby
FILE REFERENCE: 2750-1590PUS2
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT APPLICATION NUMBER: US/11/056,355B
CURRENT FILING DATE: 2004-02-14
PRIOR APPLICATION NUMBER: 60/544,190
PRIOR FILING DATE: 2004-02-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: peptide
LOCATION: (1)..(124)
OTHER INFORMATION: Ceres Seq. ID no. 12411160
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ORGANISM: Zea mays subsp. mays
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les 4; Conserv
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149 WLLAAF 154
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4; Conserva
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                                                                                                                                                                                                                         ID no. 12411159
                                                                                                                                                      Score 28; DB 7;
Pred. No. 1.3e+02;
                                                                                                                                    Mismatches
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